FINAL REPORT

STUDY TITLE

PHARMACOKINETIC (IN BLOOD)
AND EXCRETION STUDY OF [] IN RATS

Extendition brobin of [] Invited			
STUDY NUMBER			
[]			
STUDY DIRECTOR			
[]			
STUDY INITIATION DATE			
27 November 2006			
STUDY COMPLETION DATE			
26 September 2007			
PERFORMING LABORATORY			
[]			
SPONSOR			
.]			

COMPLIANCE STATEMENT

This study, designated vas conducted in compliance with the United States Environmental Protection Agency (EPA) Good Laboratory Practice (GLP) Standards (40 CFR Part 792), 18 September 1989; the Organisation for Economic Cooperation and Development (OECD) Principles of Good Laboratory Practice [C (97) 186/Final], 26 November 1997; the standard operating procedures of LLC, and the protocol as approved by the sponsor. The study was GLP compliant with the following exception. A critical phase inspection was not performed by the Quality Assurance unit while the study was in progress. A Certificate of Analysis was provided by the sponsor (presented in Appendix A); the characterization analyses were conducted according to unknown standards.

26 September 2007 Date

TABLE OF CONTENTS

		Page
	Compliance Statement	2
	Table Of Contents	3
	Index Of Tables	5
	Index Of Appendices	6
1.	Summary	7
1.1.	Objective	
1.2.	Study Design	7
1.3.	Results And Conclusions	8
2.	Introduction	9
2.1.	General Study Information	9
2.2.	Key Study Dates	9
3.	Study Design	10
4.	Experimental Procedures - Materials And Methods	11
4.1.	Test Article And Vehicle	11
.1.1.	Test Article Identification	11
.1.2.	Vehicle Identification.	11
.1.3.	Preparation	11
.1.4.	Sampling And Analyses	12
4.2.	Test System	12
4.3.	Organization Of Test Groups, Dosage Levels And Treatment Regimen	13
4.4.	Animal Receipt And Acclimation/Pretest Period	13
4.5.	Animal Housing	14
4.6.	Diet, Drinking Water And Maintenance	14
4.7.	Environmental Conditions	15
4.8.	Assignment Of Animals To Treatment Groups	15

	$\mathbf{\underline{P}_{i}}$	age
5.	Parameters Evaluated	.17
5.1.	Clinical Observations And Survival (Both Phases)	.17
5.2.	Body Weights (Both Phases)	.17
5.3.	Food Consumption (Both Phases)	.17
5.4.	Toxicokinetics	.17
5.5.	Pharmacokinetic Profile (Pharmacokinetic Phase)	.17
5.5.1.	Excretion Profile (Excretion Phase)	.17
5.6.	Study Termination (Both Phases)	.18
5.7.	Statistical Methods	.18
5.8.	Data Retention	.18
6.	Results And Discussion (Both Phases)	.19
6.1.	Analytical Chemistry	.19
6.2.	Clinical Observations And Survival	.19
6.3.	Body Weights	.19
6.4.	Toxicokinetics	
7.	Conclusions	.22
8.	Key Study Personnel And Report Submission	.23
9.	Quality Assurance Unit Statement	.24
9.1.	Phases Inspected	.24
9.2.	Approval	.26
10.	References	.27
11.	Deviations From The Protocol	.28

INDEX OF TABLES

	<u>P</u>	age
1.	Individual Survival And Disposition (Pharmacokinetic Phase)	.30
2.	Individual Survival And Disposition (Excretion Phase)	.32
3.	Individual Clinical Observations (Detailed Physical Examinations/Dispositions - Pharmacokinetic Phase)	33
4.	Individual Clinical Observations (Detailed Physical Examinations/Dispositions - Excretion Phase)	35
5.	Individual Body Weights [G] (Pharmacokinetic Phase)	.36
6.	Individual Body Weights [G] (Excretion Phase)	.38

INDEX OF APPENDICES

			<u>Pa</u>	ge
A.	Certificate Of Analysis (Sponsor-Provided Data)			40
B.	Analyses Of Dosing Formulations ([])	42
C.	Pretest Clinical Observations			88
D.	Animal Room Environmental Conditions			91
E.	Bioanalytical Report ([])		96
F.	Toxicokinetic Report ([])	2	45
G.	Study Protocol	***************************************	2	64

1. Summary

1.1. OBJECTIVE

The objectives of the study were to evaluate the pharmacokinetic (in blood) and excretion profiles of the test article in rats.

1.2. STUDY DESIGN

[] in the vehicle, sterile water for injection, was administered intravenously once to 1 pharmacokintic (blood collection) group ([]; Group 1) and 1 excretion group ([]; Group 1) of Crl:CD(SD) rats. The dosage level was 10 mg/kg for both groups at a dosage volume of 5 mL/kg. The pharmacokinetic group consisted of 9 animals/sex and the excretion group consisted of 3 animals/sex.

All animals were observed twice daily for mortality and moribundity. Detailed physical examinations were performed at least once during the pre-treatment period. Individual body weights were recorded during acclimation, at pretest initiation, at randomization and on study day 0. Food consumption was recorded during the pretest period only.

For pharmacokinetic assessment, blood samples were collected on wet ice from 3 animals/sex prior to dosing and at approximately 2, 10, 20 and 30 minutes and 1, 3, 5, 7, 24 and 48 hours after dose administration.

For excretion profile, urine samples were collected from 3 animals/sex over the following intervals: 0-6, 6-12 and 12-24 hours post-dosing.

All pharmacokinetic and excretion group animals were euthanized and discarded without further evaluation following the final blood or urine collection.

Serum and urine concentrations of [] were measured using a validated LCMS/MS method. The concentrations in serum and amounts excreted in urine were used for pharmacokinetic analysis.

1.3. RESULTS AND CONCLUSIONS

After a single intravenous dose of [] at 10 mg/kg, systemic exposure (AUC _{0-∞}) to[
for male rats was almost 7-fold higher than for female rats. [] appeared to
remain mostly in the circulation in male rats (apparent volume of distribution about
0.2 L/kg), but to have extensive tissue distribution in female rats (apparent volume of
distribution of more than 2.5 L/kg). The terminal elimination phase for [] in serum
had a half-life of 9.4 and 5.4 hours for female and male rats, respectively. The half-life
for [] in urine was 1.8 and 3.2 hours, for female and male rats respectively
Nevertheless, the percent of [] dose eliminated over 24 hours post-dosing in the urine
of male rats and female rats was similar (approximately 65%). This can be explained by
the lower amounts of [] available for urinary clearance in the circulation of female rats
compared to male rats as suggested by the differences in the apparent volume of
distribution.
In conclusion, systemic exposure (AUC $_{0-\infty}$) to [] for male rats was almost 7-fold
higher than for female rats following a single intravenous dose of [] at 10 mg/kg.
[] in the serum cleared more rapidly in the male rats than in the female rats, and the
female rats had a greater volume distribution than the male rats. [] cleared in the urine
more rapidly in the female rats than in the male rats, but most of the [] dose was
eliminated over 12 hours post-dosing in both genders. The elimination of [] in the
urine appeared to be mono-exponential for male rats and appeared not to be log-linear for
female rats

2. Introduction

2.1. GENERAL STUDY INFORMATION

This report presents the data from "Pharmacokinetic (In Blood) And Excretion Study Of

[] In Rats." Due to software spacing constraints, the study title appears as "A

Pharmacokinetic And Excretion Study In Rats" on the report tables.

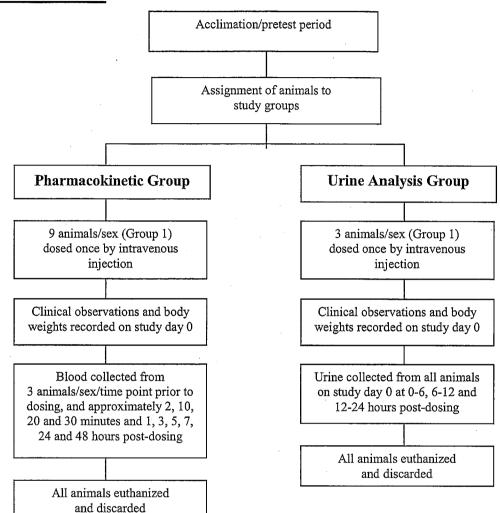
The following computer protocols were used for data collection during the study:

Computer Protocol		Type of Data Collected
[]	
[]	

2.2. KEY STUDY DATES

Date(s)	<u>Evenus)</u>
28 November 2006	Experimental starting date (animal receipt)
8 December 2006	Assignment to study groups
11 December 2006	Experimental start date (initiation of dose
	administration; study day 0)
25 July 2007	Experimental termination (completion) date
	(last bioanalytical analysis)

3. STUDY DESIGN



4. EXPERIMENTAL PROCEDURES - MATERIALS AND METHODS

4.1. TEST ARTICLE AND VEHICLE

4.1.1. TEST ARTICLE IDENTIFICATION

The test article, [
was received from Miki & Co., LTD, Chuo-Ku, Japan, on 29 November 2006, as
follows:

<u>Identification</u>	Quantity <u>Received</u>	Physical <u>Description</u>
	1 bottle	Clear, colorless
[]	Gross weight:	liquid
[log no. 7216A]	347.3 g	_

1

A Certificate of Analysis for the test article was provided by the sponsor and is presented in Appendix A. The purity of the test article was 99.0%. The test article was stored at room temperature and was considered stable under this condition. A reserve sample of the test article (0.333 g) was collected on 1 December 2006 and stored in the Archives of [______].

4.1.2. VEHICLE IDENTIFICATION

The vehicle used in preparation of the test article formulations was sterile water for injection, USP (lot no. C688010, exp. date: 1 August 2007, manufactured by Baxter Healthcare Corporation, Deerfield, Illinois).

4.1.3. PREPARATION

Dosing formulations were prepared at the concentrations indicated in the following table:

Group Number <u>Test Article</u>		Dosage Level (mg/kg)	Dosing Concentration (mg/mL) ^a	
1	[]	10	2	
^a = The dosing formulations were not adjusted for purity.				

The test article formulation was prepared as a weight/volume (test article/vehicle) mixture. The appropriate amount of the test article was weighed into a tared, labeled storage container. A predetermined volume of the vehicle was added to the container to bring the formulation nearly to the calibration mark. The formulation was mixed with a magnetic stirrer until uniform. Additional vehicle was then added to bring the formulation to the calibration mark. The formulation was stirred until uniform and stirring continued overnight in a refrigerator. The solution was removed from the refrigerator the following morning. While in a Laminar flow hood, the formulation was sterile-filtered through a 0.22-µm syringe filter into a sterile container and capped with a septum. On the day of dose administration, the formulation was removed from the refrigerator and maintained at room temperature for approximately 1 hour prior to dosing.

The test article formulations for the pharmacokinetic group (Group 1) and excretion (Group 1) group were prepared once and stored refrigerated. The test article formulation was stirred continuously throughout the preparation, sampling and dose administration procedures. A small aliquot was removed from the formulation and the pH was measured as 2.45.

4.1.4. SAMPLING AND ANALYSES

Samples (1 mL each) for concentration and stability analyses were collected from the dosing formulation at the following times: prior to filtration, after filtration and after dose administration. Homogeneity assessments were not performed as the formulations were solutions. All analyses were conducted by the Analytical Chemistry Department,

[]. The methodology and results of these analyses are presented in Appendix B, and the results are summarized in Section 6.1.

4.2. TEST SYSTEM

Crl:CD(SD) rats from Charles River Laboratories, Inc., Raleigh, North Carolina were used as the test system on this study. This species and strain of animal is recognized as appropriate for acute and subchronic toxicity studies. The Sprague-Dawley rat was used

because it is a widely used strain for which significant historical control data are available.

4.3. ORGANIZATION OF TEST GROUPS, DOSAGE LEVELS AND TREATMENT REGIMEN

The vehicle and test article formulations were administered as a single dose by a slow bolus intravenous injection using a sterile needle and syringe via a lateral tail vein. The dosage volume for all groups was 5 mL/kg. Individual doses were based on the study day 0 body weights to provide the correct mg/kg dosage.

The following tables present the study group assignment:

<u>Pharmacokii</u>	netic Group []			
Group <u>Number</u>	Test Article ^a	Dosage Level (mg/kg)	Dosage Volume (mL/kg)	Number o	of Animals <u>Females</u>
1	[]	10	5	9	9
Excretion G	coup []				
Group <u>Number</u>	Test Article ^a	Dosage Level (mg/kg)	Dosage Volume (mL/kg)	Number o	f Animals <u>Females</u>
1	[]	10	5	3	3

^a = The dosing formulations were not adjusted for purity.

The selected route of administration for this study was intravenous because this is an acceptable route of administration to assess pharmacokinetics and elimination profiles. The number of animals selected for this study was the minimum required to yield scientifically meaningful data and was consistent with regulatory agency expectations.

4.4. Animal Receipt And Acclimation/Pretest Period

Fifteen male and 15 female Crl:CD(SD) rats were received in good health on 28 November 2006, from Charles River Laboratories, Inc., Raleigh, North Carolina. The animals were approximately 38 days old at receipt. Each animal was examined by a

qualified technician on the day of receipt and weighed 3 and 10 days later. Each animal was uniquely identified by a Monel[®] metal ear tag displaying the permanent identification number. All animals were housed for at least a 7-day acclimation/pretest period. During this period, each animal was observed twice daily for mortality and changes in general appearance or behavior.

Pretest data collection began on 1 December 2006. Individual body weights were recorded and detailed physical examinations were performed periodically during the pretest period. Food consumption data were also recorded for pretest animals prior to the initiation of dose administration. Pretest clinical observations are presented in Appendix C.

4.5. Animal Housing

Upon arrival, all animals were housed individually in clean, stainless steel, wire-mesh cages suspended above cage-board. Animals were maintained in accordance with the Guide for the Care and Use of Laboratory Animals (National Research Council, 1996). The animal facilities at [] are accredited by the Association for Assessment and Accreditation of Laboratory Animal Care International (AAALAC International).

4.6. DIET, DRINKING WATER AND MAINTENANCE

The basal diet used in this study, PMI Nutrition International, LLC, Certified Rodent LabDiet® 5002, is a certified feed with appropriate analyses performed by the manufacturer and provided to []. Reverse osmosis-treated (on-site) drinking water, delivered by an automatic watering system, and the basal diet were provided ad libitum throughout the study. Municipal water supplying the facility was sampled for contaminants according to the standard operating procedures. The results of the diet and water analyses are maintained at []. No contaminants were present in animal feed or water at concentrations sufficient to interfere with the objectives of this study.

4.7. Environmental Conditions

All animals were housed throughout the acclimation period and during the dosing and sample collection phases of the study in an environmentally controlled room. The room temperature and humidity controls were set to maintain daily averages of $71\pm5^{\circ}F$ ($22\pm3^{\circ}C$) and $50\pm20\%$ relative humidity. Room temperature and relative humidity were controlled and monitored using the Metasys® DDC Electronic Environmental control system. These data were recorded approximately hourly and are summarized in Appendix D. Actual mean daily temperature ranged from $70.2^{\circ}F$ to $70.6^{\circ}F$ ($21.2^{\circ}C$ to $21.4^{\circ}C$) and mean daily relative humidity ranged from 29.3% to 45.9% during the study. Lights in the animal room were controlled by the Metasys® DDC Electronic Environmental control system and were set to provide a 12-hour light (0600 hours to 1800 hours)/12-hour dark photoperiod. The 12-hour light/12-hour dark photoperiod was interrupted as necessary to allow for the performance of protocol-specified activities. Air handling units were set to provide a minimum of 10 fresh air changes per hour.

4.8. ASSIGNMENT OF ANIMALS TO TREATMENT GROUPS

On 8 December 2006 (3 days prior to the initiation of dose administration), all available rats were weighed and examined in detail for physical abnormalities. These data were collected using the [$\,$] and reviewed by the study director. The animals judged suitable for assignment to the study were selected for use in a computerized randomization procedure. A printout containing the animal numbers, corresponding body weights and individual group assignments was generated based on body weight stratification in a block design. The animals were then arranged into groups according to the printout. Individual body weights at randomization were within $\pm 20\%$ of the mean for each sex. The pharmacokinetic group consisted of 9 males and 9 females. The excretion group consisted of 3 males and 3 females. The selected animals were approximately 7 weeks old at the initiation of dose administration; individual body weights ranged from 219 g to 248 g for males and from 157 g to 186 g

for females in the pharmacokinetic group, and from 225 g to 239 g for males and from 164 g to 174 g for females in the excretion group.

5. PARAMETERS EVALUATED

5.1. CLINICAL OBSERVATIONS AND SURVIVAL (BOTH PHASES)

All animals were observed twice daily, once in the morning and once in the afternoon, for mortality and moribundity. Detailed physical examinations were conducted approximately weekly during the pretest period.

5.2. BODY WEIGHTS (BOTH PHASES)

Individual body weights were recorded during pretest, at randomization and on study day 0.

5.3. FOOD CONSUMPTION (BOTH PHASES)

Individual food consumption was recorded during the pretest period.

5.4. TOXICOKINETICS

The methods and results of the pharmacokinetic and excretion analyses are presented in Appendix E, and the interpretation of the toxicokinetic data are presented in Appendix F.

5.5. PHARMACOKINETIC PROFILE (PHARMACOKINETIC PHASE)

Blood samples (approximately 0.5 mL each) for test article serum level determinations were collected at prior to dosing and at approximately 2, 10, 20 and 30 minutes and 1, 3, 5, 7, 24 and 48 hours after dose administration. Blood was collected via the retro-orbital sinus from isoflurane-anesthetized animals into tubes containing no anticoagulant. Samples were allowed to clot at room temperature, after which the samples were stored chilled until serum preparation. Serum was separated using a refrigerated centrifuge and frozen at approximately -20°C until transferred to the [] for analysis.

5.5.1. EXCRETION PROFILE (EXCRETION PHASE)

Urine was collected on wet ice over the following intervals: 0-6, 6-12 and 12-24 hours post-dosing. Animals were transferred to plastic metabolism cages for urine collection. At each collection interval, the cages were rinsed using a documented amount of

deionized water. Urine samples and cage rinses were frozen with minimal delay at approximately -20°C and stored at approximately -20°C until transferred to the] for analysis.

5.6. STUDY TERMINATION (BOTH PHASES)

All animals were euthanized by carbon dioxide inhalation following the final blood or urine collection and discarded without further evaluation.

5.7. STATISTICAL METHODS

Statistical analyses were not conducted on this study.

5.8. DATA RETENTION	
The sponsor has title to all documentation rec	cords, raw data, specimens or other work
product generated during the performance of	the study. All work product generated by
] , including ra	w paper data and specimens, are retained
in the Archives at [] as specified in the study protocol.
The reserve sample of the test article, pertinent	t electronic storage media and the original
final report are retained in the Archives	at [
ncompliance with regulatory requirements.	

6. RESULTS AND DISCUSSION (BOTH PHASES)

6.1. ANALYTICAL CHEMISTRY

Analytical Chemistry Report: Appendix B

The analyzed formulation was found to contain the amount of test article prescribed in the protocol (105% of target post-filter). Results of the concentration analyses of the dosing formulation are summarized below.

Text Table 3. Results of Concentration Analyses

Mean Concentration, mg/mL (% of Target), Post-Filter

Group 1

Date of Preparation

(2 mg/mL)

6, 7 December 2006

105%

The test article was found to be stable in the formulations when stored refrigerated for 5 days (102% of the corresponding time-zero value).

6.2. CLINICAL OBSERVATIONS AND SURVIVAL

Summary Data: Appendix C

Individual Data: Tables 1, 2, 3, 4

All animals survived to the scheduled euthanasia. There were no test article-related clinical observations.

6.3. BODY WEIGHTS

Individual Data: Tables 5, 6

Body weights were collected for dose calculation purposes only.

6.4. TOXICOKINETICS

Bioanalytical Report: Appendix E Toxicokinetic Report: Appendix F

The concentration of [] in the serum, urine, and cage rinse samples was measured using a validated LC-MS/MS method. The serum concentration immediately following the intravenous dose was estimated based on a regression analysis of the measured values. The mean concentrations in serum and mean amounts excreted in urine plus cage rinse were used for pharmacokinetic analysis.

The pharmacokinetic parameters for [] are summarized in the following table:

PHARMAC	OKINETIC	RESULTS F	OR [_						
		SERUM							URINE†		
10 mg/kg Intravenous Dose	C ₀ * (ng/mL)	AUC _{0-∞} (ng×h/mL)	KeI (h ⁻¹)	Half- life** (h)	Cl (L/h×kg)	V _d (L/kg)	K _{eI} (h ⁻¹)	Half- life*** (h)	% of Dose Elimi- nated		
Males	69775	373393	0.127	5.4	0.0268	0.210	0.215	3.2	67.3		
Females	102835	53137	0.074	9.4	0.188	2.55	0.392	1.8	64.0		

^{*}Values were estimated.

After a single intravenous dose of [] at 10 mg/kg, systemic exposure (AUC_{0-∞}) to [for male rats was almost 7-fold higher than for female rats. [] appeared to remain mostly in the circulation in male rats (apparent volume of distribution about 0.2 L/kg), but to have extensive tissue distribution in female rats (apparent volume of distribution of more than 2.5 L/kg). The terminal elimination phase for [] in serum had a half-life of 9.4 and 5.4 hours for female and male rats, respectively. The half-life for [] in urine was 1.8 and 3.2 hours, for female and male rats respectively. Nevertheless, the percent of [] dose eliminated over 24 hours post-dosing in the urine of male rats and female rats was similar (approximately 65%). This can be explained by the lower amounts of [] available for urinary clearance in the circulation of female rats

^{**}For the terminal elimination phase.

^{***}For urinary elimination. †Urine plus cage rinse

compared to male rats as suggested by the differences in the apparent volume of distribution.

7. Conclusions

In conclusion, systemic exposure (AUC_{0- ∞}) to [] for male rats was almost 7-fold higher than for female rats following a single intravenous dose of [] at 10 mg/kg. [] in the serum cleared more rapidly in the male rats than in the female rats, and the female rats had a greater volume distribution than the male rats. [] cleared in the urine more rapidly in the female rats than in the male rats, but most of the [] dose was eliminated over 12 hours post-dosing in both genders. The elimination of [] in the urine appeared to be mono-exponential for male rats and appeared not to be log-linear for female rats.

9. QUALITY ASSURANCE UNIT STATEMENT

9.1. PHASES INSPECTED

Date(s) of Inspection(s)	Phase Inspected	Date(s) Findings Reported to Study Director	Date(s) Findings Reported to Management	Auditor(s)
16-Feb-2007	Draft Report (Analytical Appendix)	16-Feb-2007	21-Mar-2007	M.Stauffer
12-Feb-2007 13-Feb-2007 15-Feb-2007				
16-Feb-2007	Study Records (A-1, A-2)	16-Feb-2007	21-Mar-2007	M.Stauffer
22-Feb-2007	Study Records (Rx-1)	22-Feb-2007	21-Mar-2007	K.Shaner
22-Feb-2007	Study Records (I-1)	22-Feb-2007	21-Mar-2007	K.Shaner
22-Feb-2007	Study Records (C-1)	22-Feb-2007	21-Mar-2007	K.Shaner
19-Apr-2007 20-Apr-2007 21-Apr-2007				
23-Apr-2007	Study Records (B-1, B-2, B-3)	23-Apr-2007	24-May-2007	E.Crookshank
15-May-2007	Draft Report (Bioanalytical Appendix)	15-May-2007	20-Jun-2007	E.Crookshank
06-Jun-2007 07-Jun-2007	Draft Report (Toxicokinetic Appendix)	07-Jun-2007	19-Jul-2007	E.Crookshank
06-Jun-2007 07-Jun-2007	Study Records (C-1 Supplemental, Toxicokinetic Data)	07-Jun-2007	19-Jul-2007	E.Crookshank
12-Jun-2007 13-Jun-2007	Draft Report excluding Bioanalytical Report, Toxicokinetic Report, and Analyses of Dosing Formulations			
	Appendix	13-Jun-2007	19-Jul-2007	K.Shaner
17-Sep-2007	Study Records (B-1, B-3; Long- term and freeze thaw stability)	17-Sep-2007	25-Sep-2007	E.Crookshank
18-Sep-2007	Revised Draft Report (Bioanalytical Appendix)	18-Sep-2007	25-Sep-2007	E.Crookshank

This study was inspected in accordance with the U.S. EPA Good Laboratory Practice (GLP) Standards (40 CFR Part 792), the OECD Principles of Good Laboratory Practice, the Japanese MAFF Good Laboratory Practice Standards, the standard operating

procedures of [] and the sponsor's protocol and protocol amendments, with the following exceptions. The data located in Appendix A (Certificate of Analysis) were the responsibility of the sponsor. A critical phase inspection was not performed by the Quality Assurance unit while the study was in progress. Quality Assurance findings, derived from the inspections during the conduct of the study and from the inspections of the raw data and draft report, are documented and have been reported to the study director.

This report accurately reflects the data generated during the study. The methods and procedures used in the study were those specified in the protocol, its amendments and the standard operating procedures of [

The raw data, the retention sample and the final report will be stored in the Archives at

[] or another location specified by the sponsor.

10. REFERENCES

National Research Council. *Guide for the Care and Use of Laboratory Animals,* Institute of Laboratory Animal Resources, Commission on Life Sciences; National Academy Press: Washington, DC, **1996**.

11. <u>DEVIATIONS FROM THE PROTOCOL</u>

This study was conducted in accordance with the protocol and protocol amendments, except for the following.

- **Protocol Section 6.1** states that animals will be housed 3 per cage by sex for 2 to 4 days following receipt. After consultation with the study director the animals were housed individually.
- Protocol Section 6.2 states that average daily humidity would be $50 \pm 20\%$. There were 5 or more excursions for humidity and the daily average humidity was outside acceptable range on 2 December 2006 (study day -9); the average relative humidity value for this day was 29.3%.
- **Protocol Section 7.5.2** states that analyses to demonstrate the stability of the test article formulation for the expected period of refrigerated storage between formulation and dosing will be conducted before the initiation of dosing. The dosing aliquot was inadvertently not collected until after dosing.

These deviations did not negatively impact the quality or integrity of the data nor the outcome of the study.

TABLES 1 - 6

PROJECT NO.:534006A

TABLE 1 (PHARMACOKINETIC PHASE) A PHARMACOKINETIC AND EXCRETION STUDY IN RATS INDIVIDUAL SURVIVAL AND DISPOSITION

PAGE

ANIMAL	SEX	GROUP	TYPE OF DEATH	AGE IN WEEKS A	DATE OF DEATH	DAYS ON STUDY	·
46662	M	10 MG/KG	SCHEDULED EUTHANASIA	7	13-DEC-06	2	
46663	M	10 MG/KG	SCHEDULED EUTHANASIA	7	13-DEC-06	2	
46665	M	10 MG/KG	SCHEDULED EUTHANASIA	7	13-DEC-06	2	
46666	M	10 MG/KG	SCHEDULED EUTHANASIA	7	13-DEC-06	2	
46669	M	10 MG/KG	SCHEDULED EUTHANASIA	7	13-DEC-06	2	
46672	M	10 MG/KG	SCHEDULED EUTHANASIA	7	13-DEC-06	2	
46673	M	10 MG/KG	SCHEDULED EUTHANASIA	7	13-DEC-06	2	•
46674	M	10 MG/KG	SCHEDULED EUTHANASIA	7	13-DEC-06	2	
46676	M	10 MG/KG	SCHEDULED EUTHANASIA	7	13-DEC-06	2	

A = CALCULATED TO THE NEAREST WHOLE WEEK USING THE MEAN AGE IN WEEKS AT INITIATION OF DOSING (7)

PROJECT NO.:534006A

TABLE 1 (PHARMACOKINETIC PHASE) A PHARMACOKINETIC AND EXCRETION STUDY IN RATS INDIVIDUAL SURVIVAL AND DISPOSITION

PAGE 2

							•
ANIMAL	SEX	GROUP	TYPE OF DEATH	AGE IN WEEKS A	DATE OF DEATH	DAYS ON STUDY	
46678	F	10 MG/KG	SCHEDULED EUTHANASIA	7	13-DEC-06	2	
46679	F	10 MG/KG	SCHEDULED EUTHANASIA	7	13-DEC-06	2	
46680	\mathbf{F}	10 MG/KG	SCHEDULED EUTHANASIA	7	13-DEC-06	2	•
46681	F	10 MG/KG	SCHEDULED EUTHANASIA	7	13-DEC-06	2	
46684	F	10 MG/KG	SCHEDULED EUTHANASIA	7	13-DEC-06	2	
46685	F	10 MG/KG	SCHEDULED EUTHANASIA	7	13-DEC-06	2	
46686	F	10 MG/KG	SCHEDULED EUTHANASIA	7	13-DEC-06	2	
46688	\mathbf{F}	10 MG/KG	SCHEDULED EUTHANASIA	7	13-DEC-06	2	
46691	F	10 MG/KG	SCHEDULED EUTHANASIA	7	13-DEC-06	2	

A = CALCULATED TO THE NEAREST WHOLE WEEK USING THE MEAN AGE IN WEEKS AT INITIATION OF DOSING (7)

PDEADv4.05 02/20/2007 PROJECT NO.:534006B

TABLE 2 (EXCRETION PHASE) A PHARMACOKINETIC AND EXCRETION STUDY IN RATS INDIVIDUAL SURVIVAL AND DISPOSITION

PAGE

ANIMAL	SEX	GROUP	TYPE OF DEATH	AGE IN WEEKS A	DATE OF DEATH	DAYS ON STUDY	·
46664	M	10 MG/KG	SCHEDULED EUTHANASIA	7	12-DEC-06	1	
46667	M	10 MG/KG	SCHEDULED EUTHANASIA	7	12-DEC-06	1.	
46670	M	10 MG/KG	SCHEDULED EUTHANASIA	7	12-DEC-06	1	
46682	F	10 MG/KG	SCHEDULED EUTHANASIA	7	12-DEC-06	i	
46683	F	10 MG/KG	SCHEDULED EUTHANASIA	7	12-DEC-06	1	
46690	F	10 MG/KG	SCHEDULED EUTHANASIA	7	12-DEC-06	î	

A = CALCULATED TO THE NEAREST WHOLE WEEK USING THE MEAN AGE IN WEEKS AT INITIATION OF DOSING (7)

MANUALv1.00 02/20/2007 R:02/20/2007

TABLE 3 (DETAILED PHYSICAL EXAMINATIONS/DISPOSITIONS - PHARMACOKINETIC PHASE) A PHARMACOKINETIC AND EXCRETION STUDY IN RATS

PAGE

INDIVIDUAL CLINICAL OBSERVATIONS

					TABLE 1	RANGE:	12-	11-06 TO 12-13-06
	ANIMAL S	EX	GROUP	CATEGORY	DATE	TIME	GRAD	E OBSERVATIONS
	46662		10 MG/KG	NORMAL	12-11-06			NO SIGNIFICANT CLINICAL OBSERVATIONS
		M	10 MG/KG	DISPOSITION				EUTHANIZED BY CO2 AND DISCARDED
		M	10 MG/KG	NORMAL	12-11-06			NO SIGNIFICANT CLINICAL OBSERVATIONS
		M	10 MG/KG	DISPOSITION	12-13-06	11:41	P	EUTHANIZED BY CO2 AND DISCARDED
		M	10 MG/KG	NORMAL	12-11-06			NO SIGNIFICANT CLINICAL OBSERVATIONS
		M	10 MG/KG	DISPOSITION	12-13-06			EUTHANIZED BY CO2 AND DISCARDED
		M	10 MG/KG	NORMAL	12-11-06			NO SIGNIFICANT CLINICAL OBSERVATIONS
		M	10 MG/KG	DISPOSITION				EUTHANIZED BY CO2 AND DISCARDED
		M	10 MG/KG	NORMAL	12-11-06			NO SIGNIFICANT CLINICAL OBSERVATIONS
		M	10 MG/KG	DISPOSITION	12-13-06			EUTHANIZED BY CO2 AND DISCARDED
		M	10 MG/KG	NORMAL	12 -11 -06	8:11	P	NO SIGNIFICANT CLINICAL OBSERVATIONS
		M	10 MG/KG	DISPOSITION	12-13-06			EUTHANIZED BY CO2 AND DISCARDED
		M	10 MG/KG	NORMAL	12-11-06			NO SIGNIFICANT CLINICAL OBSERVATIONS
		M	10 MG/KG	DISPOSITION	12-13-06			EUTHANIZED BY CO2 AND DISCARDED
		M	10 MG/KG	NORMAL	12-11-06	8:11	P	NO SIGNIFICANT CLINICAL OBSERVATIONS
		M	10 MG/KG	DISPOSITION	12-13-06			EUTHANIZED BY CO2 AND DISCARDED
.)		M	10 MG/KG	NORMAL	12-11-06			NO SIGNIFICANT CLINICAL OBSERVATIONS
		M	10 MG/KG	DISPOSITION	12-13-06	11:42	P	EUTHANIZED BY CO2 AND DISCARDED
,		F	10 MG/KG	NORMAL	12-11-06			NO SIGNIFICANT CLINICAL OBSERVATIONS
š		F	10 MG/KG	DISPOSITION	12-13-06			EUTHANIZED BY CO2 AND DISCARDED
ō		F	10 MG/KG	NORMAL	12-11-06			NO SIGNIFICANT CLINICAL OBSERVATIONS
_		F	10 MG/KG	DISPOSITION	12-13-06	11:42	P	EUTHANIZED BY CO2 AND DISCARDED
		F	10 MG/KG	NORMAL	12-11-06	8:12	P	NO SIGNIFICANT CLINICAL OBSERVATIONS
		F	10 MG/KG	DISPOSITION	12-13-06	11:42	\mathbf{P}	EUTHANIZED BY CO2 AND DISCARDED
		F	10 MG/KG	NORMAL	12-11-06	8:13	P	NO SIGNIFICANT CLINICAL OBSERVATIONS
		F	10 MG/KG	DISPOSITION	12-13-06		P	EUTHANIZED BY CO2 AND DISCARDED
		F	10 MG/KG	NORMAL	12-11-06			
	46684		10 MG/KG	DISPOSITION	12-13-06	11:42		EUTHANIZED BY CO2 AND DISCARDED
	46685	F	10 MG/KG	NORMAL	12-11-06	8:13	P	NO SIGNIFICANT CLINICAL OBSERVATIONS

GRADE CODE: 1 - SLIGHT 2 - MODERATE 3 - SEVERE P - PRESENT

33 of 281

PROJECT NO.:534006A

TABLE 3 (DETAILED PHYSICAL EXAMINATIONS/DISPOSITIONS - PHARMACOKINETIC PHASE)
A PHARMACOKINETIC AND EXCRETION STUDY IN RATS
INDIVIDUAL CLINICAL OBSERVATIONS

TABLE RANGE: 12-11-06 TO 12-13-06

ANIMAL SEX	GROUP	CATEGORY	DATE TIME GRADE OBSERVATIONS
46685 F 46686 F 46686 F 46688 F 46688 F 46691 F 46691 F	10 MG/KG 10 MG/KG 10 MG/KG 10 MG/KG 10 MG/KG 10 MG/KG 10 MG/KG	DISPOSITION NORMAL DISPOSITION NORMAL DISPOSITION NORMAL DISPOSITION	12-13-06 11:42 P EUTHANIZED BY CO2 AND DISCARDED 12-11-06 8:13 P NO SIGNIFICANT CLINICAL OBSERVATIONS 12-13-06 11:42 P EUTHANIZED BY CO2 AND DISCARDED 12-11-06 8:14 P NO SIGNIFICANT CLINICAL OBSERVATIONS 12-13-06 11:42 P EUTHANIZED BY CO2 AND DISCARDED 12-11-06 8:14 P NO SIGNIFICANT CLINICAL OBSERVATIONS 12-13-06 11:42 P EUTHANIZED BY CO2 AND DISCARDED

GRADE CODE: 1 - SLIGHT 2 - MODERATE 3 - SEVERE P - PRESENT

PCRDv4.11 02/20/2007 R:02/21/2007

PAGE

PROJECT NO.:534006A

TABLE 4 (DETAILED PHYSICAL EXAMINATIONS/DISPOSITIONS - EXCRETION PHASE)
A PHARMACOKINETIC AND EXCRETION STUDY IN RATS
INDIVIDUAL CLINICAL OBSERVATIONS

TABLE RANGE: 12-11-06 TO 12-12-06

ANIMAL	SEX	GROUP	CATEGORY	DATE 1	TIME GRA	ADE OBSERVATIONS		
46664 46667 46667 46670 46670 46682 46682 46683 46683 46690	M M M M F F F F	10 MG/KG 10 MG/KG	NORMAL DISPOSITION	12-11-06 12-12-06 12-11-06 12-11-06 12-11-06 12-11-06 12-11-06 12-11-06 12-11-06 12-11-06 12-11-06	12:00 P 8:17 P 12:01 P 8:17 P 12:01 P 8:17 P 12:01 P 8:17 P 12:01 P 8:18 P	P BUTHANIZED BY CO2 AND DISCARDED P NO SIGNIFICANT CLINICAL OBSERVATIONS P EUTHANIZED BY CO2 AND DISCARDED P NO SIGNIFICANT CLINICAL OBSERVATIONS P EUTHANIZED BY CO2 AND DISCARDED P NO SIGNIFICANT CLINICAL OBSERVATIONS P EUTHANIZED BY CO2 AND DISCARDED P NO SIGNIFICANT CLINICAL OBSERVATIONS P EUTHANIZED BY CO2 AND DISCARDED P NO SIGNIFICANT CLINICAL OBSERVATIONS P EUTHANIZED BY CO2 AND DISCARDED P NO SIGNIFICANT CLINICAL OBSERVATIONS		

GRADE CODE: 1 - SLIGHT 2 - MODERATE 3 - SEVERE P - PRESENT

PCRDv4.11 02/20/2007 R:02/21/2007

PAGE

PROJECT NO.:534006B

PROJECT NO.:534006A

TABLE 5 (PHARMACOKINETIC PHASE) A PHARMACOKINETIC AND EXCRETION STUDY IN RATS INDIVIDUAL BODY WEIGHTS [G]

PAGE

DAY	-10	-3	· O	MALE GROUP:	10 MG/KG
ANIMAL					
46662	152.	218.	241.		
46663	153.	227.	248.		
46665	142.	206.	226.		
46666	138.	201.	223.		
46669	145.	210.	236.		
46672	147.	221.	242.		•
46673	137.	207.	233.	•	
46674	150.	207.	228.		
46676	141.	201.	219.		
MEAN	145.	211.	233.		
S.D.	5.9	9.1	9.7		
N	9	9	9		

PROJECT NO.:534006A

TABLE 5 (PHARMACOKINETIC PHASE) A PHARMACOKINETIC AND EXCRETION STUDY IN RATS INDIVIDUAL BODY WEIGHTS [G]

PAGE

DAY	-10	-3	0	FEMALE GROUP:	10 MG/KG
ANIMAL 46678	120.	155.	162.		
46679 46680	139. 138.	174. 163.	186. 166.		
46681 46684	107. 131.	162. 170.	162. 177.		
46685 46686	122. 140.	151. 175.	157. 186.		
46688 46691	126. 117.	157. 162.	163. 167.		
MEAN S.D.	127. 11.3	163. 8.4	170. 10.8		
N.	9	9	9		

PBFTSv4.44 02/20/2007

37 of 28

PROJECT NO.:534006B

TABLE 6 (EXCRETION PHASE) A PHARMACOKINETIC AND EXCRETION STUDY IN RATS INDIVIDUAL BODY WEIGHTS [G]

PAGE

DAY	-10	-3	0	MALE GROUP: 10 MG/KG
ANTMAL 46664 46667 46670	144. 147. 146.	206. 216. 213.	225. 239. 235.	·
MEAN S.D. N	146. 1.5 3	212. 5.1 3	233. 7.2 3	

PROJECT NO.:534006B

TABLE 6 (EXCRETION PHASE) A PHARMACOKINETIC AND EXCRETION STUDY IN RATS INDIVIDUAL BODY WEIGHTS [G]

PAGE

DAY	-10	-3	0	FEMALE GROUP: 10 MG/KG
ANIMAL 46682 46683 46690	135. 136. 140.	167. 164. 161.	168. 174. 164.	
MEAN S.D. N	137. 2.6 3	164. 3.0 3	169. 5.0 3	PBFTSv4.44 02/20/2007

39 of 28

APPENDIX A

Certificate of Analysis (Sponsor Provided Data)

Item	Unit	Analysis Results	
	AREA%	99.62	
			·
unkoown	. AREA%	0.005	
unknown		0. 005	
unknown		0. 058	
unknown		0. 010	·
unknown		0. 029	
unknown		0. 183	
unknown .		0.024	
unknown		0. 035	
unknown .		0.027	
			·
		•	
			•
```			

(This is stable for 3 years in room temperature.)

Note: The issue of this analysis shall be recognized as authorized values herein given by the Manager, Quality Assurance Office.

### APPENDIX B

Analyses Of Dosing Formulations [

Analyses Of Dosing Formulations

]

-1-

43 of 281

## TABLE OF CONTENTS

	<u>Page</u>
	Table Of Contents2
	Index Of Figures3
	Index Of Tables
	Index Of Attachments
1.	Summary4
2.	Introduction5
3.	Experimental5
3.1.	Instrument5
3.2.	High Performance Liquid Chromatography5
3.3.	Mass Spectrometry6
3.4.	Preparation Of Mobile Phase/Diluent
3.5.	Preparation Of The Calibration Standard Stock Solution And Calibration Standards
3.6.	Preparation Of Quality Control Stock Solution
3.7.	Preparation Of Quality Control Samples
3.8.	Sample Processing9
3.9.	Calibration And Quantitation9
4.	Results And Discussion
4.1.	Specificity/Selectivity
4.2.	Assay Validation: Calibration Reproducibility
4.3.	Assay Validation: Precision And Accuracy
4.4.	Stability Of Processed Samples
4.5.	Concentration Analysis Of Dosing Formulations
4.6.	Test Article Stability In Formulations
5.	Conclusion14
6.	Key Study Personnel And Report Submission16

### INDEX OF FIGURES

	Page
Figure 1:	Representative Chromatogram Of A 250 ng [ ]/mL Calibration Standard
Figure 2:	Representative Chromatogram Of A Processed 2.00 mg [ ]/mL Quality Control Sample
Figure 3:	Representative Chromatogram Of A Processed 2 mg [ ]/mL Formulation Sample
Figure 4:	Representative Chromatogram Of A Processed Vehicle Sample12
	INDEX OF TABLES
Table 1:	[ ] Back-Calculated Concentrations And Intra-Session Statistics Of Calibration Standards
Table 2:	Control Samples20
Table 3:	Room Temperature Stability Of [ ] In Processed Samples22
Table 4:	Concentration Analysis Of The 6 And 7 December 2006 Formulation
Table 5:	5-Day Refrigerated Stability Analysis Of The 6 And 7 December 2006 Formulation
	INDEX OF ATTACHMENTS
I:	Supporting Data

#### 1. SUMMARY

This report provides a detailed description of a high performance liquid chromatography tandem mass spectrometry method in the negative Turbo ionspray mode for the determination of [

[ ] concentration in sterile water. Method specificity/selectivity, calibration reproducibility, precision and accuracy were assessed and validated.

Quantitation was performed using calibration standards in the range of 250 to 3000 ng [ ]/mL. The inter-set variability of the back-calculated standard concentrations ranged from 0.95% to 2.4% relative standard deviation (RSD). The inter-set mean standard concentrations had percent relative error (%RE) values ranging from -1.4% to 1.1%.

Precision and accuracy were verified by the analysis of quality control (QC) samples prepared at 2.00 mg [ ]/mL in sterile water. The inter-set variability of the calculated QC concentration (precision) was 5.2% RSD. The inter-session mean QC concentration had a %RE value (accuracy) of 4.2%.

The formulation used for dosing was prepared at an [ ] concentration of 2 mg/mL and was assessed for test article concentration prior to and after filtration. The post-filtration results met the standard operating procedure (SOP) requirements, i.e., the concentration was within the acceptable limits (within 90% to 110% of target concentration).

In addition, test article stability in the formulation prepared at 2 mg [ ]/mL was evaluated for up to 5 days of refrigerated storage. The test article in the formulation met the [ ] SOP requirement for stability, i.e., the mean post-storage analyte concentration was not less than 90% of the time-zero concentration.

#### 2. Introduction

This report provides a detailed description of a high performance liquid chromatography tandem mass spectrometry (HPLC/MS/MS) method in the negative Turbo ionspray mode for the determination of [

]concentration in sterile water. Method specificity/selectivity, ruggedness, calibration reproducibility, precision and accuracy were assessed and validated. In addition, dosing formulations were analyzed to confirm test article concentration and stability following refrigerated storage for 5 days.

#### 3. EXPERIMENTAL

#### 3.1. Instrument

A Spark Holland, Inc. Symbiosis Pharma System was used for the HPLC analysis. The Symbiosis Pharma consisted of a gradient pump set, a Reliance autosampler with Conditioned Stacker and a Spark Holland Mistral column heater. An Applied Biosystems, Inc. API 5000 tandem mass spectrometer equipped with TurboIonSpray was used for the mass monitoring. Data acquisition and analysis were performed using Analyst[®] 1.4.1. (Note: the retention and run time may have varied depending on column performance. Mass spectrometer conditions may have varied depending on mass spectrometer performance).

#### 3.2. HIGH PERFORMANCE LIQUID CHROMATOGRAPHY

Column:

ACE C8  $50 \times 2.1$  mm with a C8 guard cartridge, or

equivalent

Column Temperature:

28°C to 30°C

Mobile Phase:

2 mM ammonium acetate in 70:30 deionized

water:acetonitrile

Flow Rate:

0.3 mL/minute

Detector:

Mass spectrometer with conditions as described in

Section 3.3. (Mass Spectrometry)

Injection Volume:

 $5 \mu L$ 

Retention Time:

Approximately 3.2 minutes for [ ]

Run Time:

5 minutes

### 3.3. MASS SPECTROMETRY

### **Acquisition Parameters**

Scan Type:

Multiple reaction monitoring (MRM)

Polarity:

Negative

Scan Mode:

N/A

Ion Source:

Turbo Spray

Resolution Q1:

Unit

Resolution Q3:

Unit

Intensity Threshold:

0.00 cps

Settling Time:

0.00 msec

MR Pause:

5.0070 msec

MCA:

No

Step Size:

0.00 amu

<u>Analyte</u>	<u>Q1 Mass</u> (amu)	Q3 Mass (amu)	<u>Dwell</u> (msec)	<u>Parameter</u>	Start/Stop
[]	[ ]	[ ]	[ ]	DP	30/-30
				CE	-11/-11
				CXP	-4.0/-4.0

DP = Declustering potential

CE = Collision energy

CXP = Collision cell exit potential

#### Parameter Table

CUR (Curtain Gas): 10.00

GS1 (Gas 1): 70.00

GS2 (Gas 2): 20.00

IS (IonSpray Voltage): -4000.00

TEM (Temperature): 450.00

ihe (Interface Heater): ON

CAD (Collisionally Activated

Dissociation Gas):

4.00 (nitrogen)

EP (Entrance Potential): -12.00

#### **Detection Parameters (Negative)**

CEM (Channel Electron Multiplier): 2600.0

DF (Deflector): 200.0

### 3.4. Preparation Of Mobile Phase/Diluent

The mobile phase/diluent (2 mM ammonium acetate in 70:30 deionized water:acetonitrile) was prepared by dissolving approximately 0.154 g of ammonium acetate in 700 mL of deionized (DI) water. Acetonitrile (ACN, 300 mL) was added, and the solution was stirred to achieve complete dissolution and vacuum-degassed. The preparation was scaled as needed.

# 3.5. PREPARATION OF THE CALIBRATION STANDARD STOCK SOLUTION AND CALIBRATION STANDARDS

A calibration standard stock solution was prepared at a concentration of 200  $\mu$ g [ ]/mL as follows. Approximately 0.0100 g of [ ] ([ ] log no. 7216A) was accurately weighed in a tared glass weigh funnel and transferred to a 50-mL volumetric flask with rinses of DI water. Additional DI water was added as needed to yield the desired concentration, and the solution was stirred to achieve complete dissolution.

An aliquot (0.5 mL) of the calibration standard stock solution was transferred to a 15-mL polypropylene tube containing 9.5 mL of mobile phase to yield a working stock solution at a concentration of 10.0  $\mu$ g [ ]/mL. Dilutions of this working stock solution were prepared in mobile phase in autosampler vials to yield calibration standards spanning the concentration range of 250 to 3000 ng [ ]/mL.

#### 3.6. Preparation Of Quality Control Stock Solution

A quality control (QC) stock solution was prepared at a concentration of 5.0 mg [ ]/mL as follows. Approximately 0.126 g of [ ] log no. 7216A) was accurately weighed in a tared 25-mL volumetric flask. DI water was added as needed to yield the desired concentration, and the solution was stirred to achieve complete dissolution.

#### 3.7. PREPARATION OF QUALITY CONTROL SAMPLES

The QC samples were prepared in triplicate. Aliquots of the QC stock solution were added to 1.0 mL of sterile water in 50-mL polypropylene tubes to achieve QC sample concentrations of 2.0 mg/mL. Appropriate volumes of mobile phase were added to each tube to achieve a final volume of 40.0 mL, and the samples were thoroughly mixed. Secondary dilutions were prepared with mobile phase in 5.0-mL polypropylene tubes. A portion of each QC sample was transferred to an amber autosampler vial for analysis.

	Initial	Sterile	Stock QC	Diluent	Total	Diluted
<u>Level</u>	<u>Concentration</u>	<u>Water</u>	<u>Volume</u>	<u>Volume</u>	<u>Volume</u>	Concentration
	(mg/mL)	(mL)	(mL)	(mL)	(mL)	$(\mu g/mL)$
Blank	0.0	1.0	0.0	39.0	40.0	0.0
QC	2.0	1.0	0.400	38.6	40	50.0

#### **Secondary Dilutions**

<u>Level</u>	Initial Concentration (mg/mL)	Aliquot <u>Volume</u> (mL)	Diluent Volume (mL)	Total <u>Volume</u> (mL)	Final Concentration (ng/mL)
Blank	0.0	0.080	3.92	4.0	0.0
QC	2.0	0.080	3.92	4.0	1000

#### 3.8. SAMPLE PROCESSING

Formulation samples (1.0 mL) were diluted with 39.0 mL of mobile phase in 50-mL polypropylene tubes. The samples were thoroughly mixed, and a secondary dilution was prepared with mobile phase in new polypropylene tubes. A portion of each sample was transferred to an amber autosampler vial for analysis.

	Test Article	Sample	Diluent	Total	Diluted
<u>Group</u>	<b>Concentration</b>	<u>Volume</u>	<u>Volume</u>	<u>Volume</u>	Concentration
	(mg/mL)	(mL)	(mL)	(mL)	$(\mu g/mL)$
1	2	1.0	39.0	40.0	50

#### **Secondary Dilutions**

Group	Test Article Concentration (mg/mL)	Aliquot <u>Volume</u> (mL)	Diluent Volume (mL)	Total Volume (mL)	Final Concentration (ng/mL)
1	2	0.080	3.92	4.0	1000

### 3.9. CALIBRATION AND QUANTITATION

Single injections were made of each calibration standard and processed QC and formulation sample. A calibration curve was constructed for each set of analyses using Analyst[®] software. The [ ] peak areas (y) and the theoretical concentrations of the calibration standards (x) were fit to the quadratic function with  $1/x^2$  weighting, excluding the origin. Concentrations were calculated using Analyst[®] software. The concentration

data were transferred to an Excel spreadsheet, where appropriate summary statistics, i.e., mean, standard deviation (SD), relative standard deviation (RSD), percent relative error (%RE) and percent of target, were calculated and presented in tabular form. The concentrations of the dosing formulations and QC samples were calculated by applying any necessary multiplication factors

#### 4. RESULTS AND DISCUSSION

Under the described chromatographic conditions, the retention time of the test article was approximately 3.2 minutes. Figures 1, 2, 3 and 4 are typical chromatograms of a calibration standard, a processed QC sample, a processed formulation sample and a processed vehicle sample, respectively. The total analysis time required for each run was 5 minutes.

#### 4.1. SPECIFICITY/SELECTIVITY

As shown in Figure 4 (and in contrast to the chromatograms shown in Figures 1 through 3), assay specificity/selectivity was confirmed when HPLC/MS/MS analysis of a processed vehicle sample revealed that there were no significant peaks at or near the retention time for the test article (approximately 3.2 minutes).

### 4.2. ASSAY VALIDATION: CALIBRATION REPRODUCIBILITY

During each of 4 validation sessions, triplicate calibration standards at 5 concentrations were prepared and analyzed as previously described. Single injections were made of each calibration standard. The resulting peak area versus concentration data were fit to the quadratic function with  $1/x^2$  weighting, excluding the origin, using least-squares regression analysis. The results of the regression analyses were used to back-calculate the corresponding concentrations from the peak area data. The reproducibility of the calibration curve data was considered valid when 1) the inter-session variability of the back-calculated concentrations at each calibration level was  $\leq 10\%$  RSD, except at the lowest calibration level where  $\leq 15\%$  RSD was acceptable; and 2) the mean back-calculated concentrations at each calibration level were within 10% of the

theoretical values (%RE within  $\pm$  10%), except at the lowest calibration level where %RE within  $\pm$  15% was acceptable.

The back-calculated concentrations and the associated intra- and inter-session statistics for the [ ] assay calibration standards are summarized in Table 1. The inter-session variability of the back-calculated concentrations ranged from 0.95% to 2.4% RSD. The inter-session mean concentrations had %RE values ranging from -1.4% to 1.1%. Based on the stated criteria, the reproducibility of the calibration data was acceptable.

#### 4.3. ASSAY VALIDATION: PRECISION AND ACCURACY

During each of 4 validation sessions, triplicate QC samples at a single concentration were prepared and analyzed as described previously. Single injections were made of each processed QC sample. The results of the regression analyses were used to back-calculate the corresponding concentrations from the QC peak area data. The variability (RSD) of the calculated QC concentration data was used as a measure of assay precision. The precision of the method was considered acceptable when the inter-session RSD of the calculated QC concentration was  $\leq 10\%$ . The difference between the theoretical and mean calculated QC concentration (%RE) was used as a measure of assay accuracy. The accuracy of the method was considered acceptable when the inter-session mean calculated concentration had a %RE value within  $\pm 10\%$ .

The calculated concentrations and the associated intra- and inter-session statistics for the calculated concentration was 5.2% RSD. The inter-session mean concentration had a %RE value of 4.2%.

1

Based on the stated criteria, the precision and accuracy of the [ ] assay were acceptable.

#### 4.4. STABILITY OF PROCESSED SAMPLES

Calibration standards and processed QC samples were analyzed. The 250 and 3000 ng/mL calibration standards and the processed QC samples were then stored at room temperature for 1 day and reanalyzed to assess test article stability. The mean test article concentrations in the calibration standards and processed QC samples after storage ranged from 102% to 105% of the time-zero values (Table 3), which met the [ ] standard operating procedure (SOP) requirement for stability, i.e., the mean post-storage concentration was not less than 90% of the time-zero value.

#### 4.5. CONCENTRATION ANALYSIS OF DOSING FORMULATIONS

The results of the determination of the test article concentration in a sterile water formulation prepared on 6 and 7 December 2006 are presented in Table 4. The concentration prior to filtration was 2.07 mg/mL (103% of target). The mean post-filtration concentration was 2.11 mg/mL (105% of target). The analyzed post-filtration [ ] formulation used for dose administration met the [ ] SOP requirement for concentration acceptability for solution formulations, i.e., the analyzed concentrations were within 90% to 110% of the target concentrations.

### 4.6. TEST ARTICLE STABILITY IN FORMULATIONS

The formulation prepared on 6 December 2006 and initially analyzed on 7 December 2006 was refrigerated for 5 days and reanalyzed to assess test article stability in the formulation. The 5-day stability results for the formulation are presented in Table 5. The mean post-storage concentration was 102% of the corresponding time-zero value, which met the previously stated [ ] SOP requirement for stability.

#### 5. CONCLUSION

This report provides a detailed description of an HPLC/MS/MS method in ESI- mode for the determination of [ ] in sterile water. Method specificity/selectivity, calibration reproducibility, precision and accuracy were assessed and validated. The analyzed [ ] formulations were analyzed to assess test article concentration prior to and after filtration,

and the results met all appropriate [ ] SOP requirements. Test article stability in formulation stored refrigerated for 5 days was assessed, and the results met [ ] SOP acceptance criteria. The analyzed [ ] formulation used for dose administration met the [ ] SOP requirement for concentration acceptability for solution formulations.

			·	
	·			
·				

# TABLES 1-5

Table 1: [ ] Back-Calculated Concentrations And Intra-Session Statistics Of Calibration Standards

Theo. Conc. (ng/mL)	250	750	1000	2000	3000
Set 1 (5-6 Dec 2006), I6-53	4006d, ana	lyst DKP		<u> </u>	
Sample 1	252	752	994	1976	2999
Sample 2	252	751	1007	1971	2996
Sample 3	245	748	1013	1992	3056
Intra-set Statistics					
n	3	3	3	3	3
Mean	250	750	1005	1980	3017
SD	4.3	2.5	9.5	11	34
RSD	1.7	0.33	0.95	0.55	1.1
%RE	-0.056	0.063	0.47	-1.0	0.56
Set 2 (5-6 Dec 2006), I6-53	4006e, anal	yst DKP			
Sample 1	253	757	1012	1912	2970
Sample 2	247	739	1036	1936	3052
Sample 3	250	743	996	2037	3068
Intra-set Statistics					
n	3	3	3	3	3
Mean	250	747	1015	1962	3030
SD	2.9	9.5	20	66	53
RSD	1.2	1.3	2.0	3.4 `	1.7
%RE	-0.060	-0.45	1.5	-1.9	0.99
Set 3 (7 Dec 2006), 16-5340	06f, analysi	t DKP			
Sample 1	256	750	998	1953	3008
Sample 2	242	730	1019	1962	3027
Sample 3	252	757	1017	2020	3016
Intra-set Statistics					
n	3	3	3	3	3
Mean	250	746	1011	1978	3017
SD	7.3	14	12	36	9.5
RSD	2.9	1.9	1.1	1.8	0.32
%RE	-0.0027	-0.56	1.1	-1.1	0.56

Table 1: [ ] Back-Calculated Concentrations And Intra-Session
Statistics Of Calibration Standards

Theo. Conc. (ng/mL)	250	750	1000	2000	3000					
Set 4 (11 Dec 2006), I6-534006g, analyst DKP										
Sample 1	259	726	998	1937	3039					
Sample 2	252	737	1022	1943	3006					
Sample 3	238	778	1014	2035	3019					
Intra-set Statistics										
n	3	3	3	3	3					
Mean	250	747	1011	1972	3021					
SD	11	27	12	55	17					
RSD	4.3	3.7	1.2	2.8	0.56					
%RE	-0.040	-0.37	1.1	-1.4	0.71					
Inter-set Statistics										
n	12	12	12	12	12					
Mean	250	748	1011	1973	3021					
SD	5.9	14	13	41	29					
RSD	2.4	1.9	1.2	2.1	0.95					
%RE	-0.040	-0.33	1.1	-1.4	0.70					

Table 2: [ ] Concentrations And Intra-Session Statistics Of Quality Control Samples

Theo. Conc. (mg/mL)	2.00				
Set 1 (5-6 Dec 2006), I6-5340	06d, analyst DKP				
Sample 1	2.00				
Sample 2	2.04				
Sample 3	2.03				
Intra-set Statistics					
n	3				
Mean	2.02				
SD	0.024				
RSD	1.2				
%RE	1.2				
Set 2 (5-6 Dec 2006), I6-5340	06e, analyst DKP				
Sample 1	2.24				
Sample 2	2.28				
Sample 3	2.27				
Intra-set Statistics					
n	3				
Mean	2.26				
SD	0.024				
RSD	1.1				
%RE	13				
Set 3 (7 Dec 2006), 16-534006	f, analyst DKP				
Sample 1	2.03				
Sample 2	2.04				
Sample 3	2.02				
Intra-set Statistics					
n	3				
Mean	2.03				
SD	0.0094				
RSD	0.46				
%RE	1.5				

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Table 3: Room Temperature Stability Of [ ] In Processed Samples

Storage <u>Duration</u> (Days)	Theo. Conc. (ng/mL)	<u>Run #</u> (534006-)	<u>Ref. #</u> (534006-)	Analyzed <u>Conc.</u> (ng/mL)	% Target	Mean <u>Conc</u> (ng/mL)	RSD (%)	Mean <u>% Target</u>	% of Time Zero
0	250	69	8-2	252	101	250	1.7	99.9	NA
		70	8-3	252	101				
		71	8-4	245	98.0				
	3000	81	8-14	2999	100	3017	1.1	101	NA
		82	8-15	2996	99.9				
		83	8-16	3056	102				
1	250	138	8-2	260	104	261	0.83	104	104
		139	8-3	263	105				
		140	8-4	259	104				
	3000	144	8-14	3177	106	3177	1.3	106	105
		145	8-15	3137	105				
		146	8-16	3217	107				
Storage	Theo.			Analyzed	%	Mean		Mean	% of Time
Duration (Days)	Conc. (mg/mL)	<u>Run #</u> (534006-)	<u>Ref. #</u> (534006-)	Conc. (mg/mL)	Target	Conc (mg/mL)	RSD (%)	% Target	<u>Zero</u>
0	2.0	85	9-7	2.00	99.8	2.02	1.2	101	NA
		86	9-8	2.04	102				
		87	9-9	2.03	102				
1	2.0	141	9-7	2.04	102	2.06	1.1	103	102
		142	9-8	2.08	104				
		143	9-9	2.07	103	•			
					÷		53400	6 report tabl	es.xls PSS

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#### Table 4: Concentration Analysis Of The 6 And 7 December 2006 Formulation

(Analyzed 07 December 2006)

				(	· · · · · · · · · · · · · ·				
Dose				Analyzed	Percent of	Mean			Mean Conc
<u>Conc</u>	<u>Group</u>	<u>Ref #</u>	<u>Run #</u>	<u>Conc</u>	<u>Target</u>	Conc	<u>SD</u>	<u>RSD</u>	% of Target
(mg/mL)		(534006-)	(534006)	( mg/mL )	(%)	(mg/mL)		(%)	(%)
Concentration	of dosing pre	parations prior	to filtration						
_		•							
2	1	17 - 5	133	2.44	122	2.25	0.27	12	113
		17 - 6	134	2.07	103				
Concentration	of dosing pre	eparations after	filtration						
2	1	17 - 7	135	2.09	105	2.11	0.021	0.99	105
		17 - 8	136	2.12	106				

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#### Table 5: 5-Day Refrigerated Stability Analysis Of The 6 And 7 December 2006 Formulation (Analyzed 11-12 Dec 2006)

Group	Dose Conc. ( mg/mL )	<u>Ref#</u> (534006 - )	<u>Run #</u>	Analyzed Conc. (mg/mL)	Percent of <u>Target</u> (%)	Mean Conc. ( mg/mL )	<u>SD</u>	<u>RSD</u> (%)	Mean Conc % of Target (%)	Percent of Time Zero (%)
1	2 -	24 - 3 24 - 4	173 174	2.16 2.13	108 106	2.15	0.025	1.2	107	102

(mg/mL) Time Zero Concentration:

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### ATTACHMENT I

Supporting Data

Table A-1: 16-534006d Data

Note: Validation Session I QC samples are listed as the diluted concentration of QC 1000.

-26-

68 of 281

	File Name	Sample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record	Analyte Concentration (ng/mL)	Dilution Factor
	l6-534006d\0066.wiff	8-2	system suit	Unknown	V-1/2		N/A	1.00
	I6-534006d\0067.wiff	8-2	system suit	Unknown	***************************************	······································	N/A	1.00
	I6-534006d\0068.wiff	8-2	system suit	Unknown	***************************************		N/A	1.00
	I6-534006d\0069.wiff	8-2	C 250	Standard	·····	×	250.00	1.00
	I6-534006d\0070.wiff	8-3	C 250	Standard			250.00	1.00
	I6-534006d\0071.wiff	8-4	C 250	Standard			250.00	1.00
	I6-534006d\0072.wiff	8-5	C 750	Standard		~~~~	750.00	1.00
	l6-534006d\0073.wiff	8-6	C 750	Standard	***************************************	***************************************	750.00	1.00
	I6-534006d\0074.wiff	8-7	C 750	Standard	-	$\square$	750.00	1.00
	I6-534006d\0075.wiff	8-8	C 1000	Standard	·····	×	1000.0	1.00
	I6-534006d\0076.wiff	8-9	C 1000	Standard		$\boxtimes$	1000.0	1.00
	I6-534006d\0077.wiff	8-10	C 1000	Standard		×	1000.0	1.00
45	I6-534006d\0078.wiff	8-11	C 2000	Standard	***************************************		2000.0	1.00
	l6-534006d\0079.wiff	8-12	C 2000	Standard	***************************************		2000.0	1.00
	I6-534006d\0080.wiff	8-13	C 2000	Standard			2000.0	1.00
3	I6-534006d\0081.wiff	8-14	C 3000	Standard			3000.0	1.00
	I6-534006d\0082.wiff	8-15	C 3000	Standard	,		3000.0	1.00
	l6-534006d\0083.wiff	8-16	C 3000	Standard		$\boxtimes$	3000.0	1.00
Ú,	l6-534006d\0084.wiff	9-6	QC 0	Quality Control		$\boxtimes$	0.0000	1.00
	16-534006d\0085.wiff	9-7	QC 1000	Quality Control			1000.0	1.00
	l6-534006d\0086.wiff	9-8	QC 1000	Quality Control			1000.0	1.00
	I6-534006d\0087.wiff	9-9	QC 1000	Quality Control		*************************	1000.0	1.00

Printing Date: Thursday, December 07, 2006 Printing Time: 9:17:59 AM

Operator: ullman Analyst Version: 1.4.1

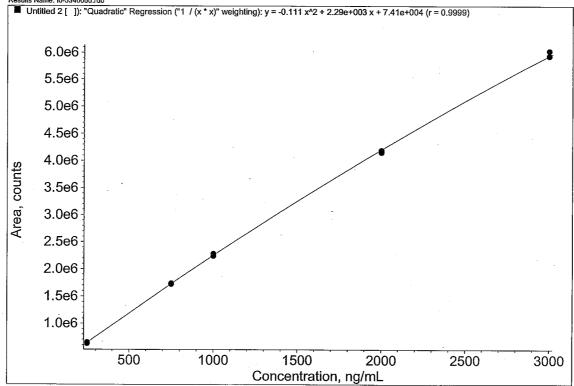
	File Name	Calculated Concentration (ng/mL)	%RE	Analyte Peak Area (counts)	Analyte Retention Time (min)	Analyte Integration Type	Record Modified	Sample Annotation
1	I6-534006d\0066.wiff	251.19	N/A	6.42e+005	3.13	Base To Base	1 - 1	
2	I6-534006d\0067.wiff	243.65	N/A	6.25e+005	3.18	Base To Base		<u> </u>
3	I6-534006d\0068.wiff	247.06	N/A	6.32e+005	3.17	Base To Base		
4	I6-534006d\0069.wiff	252.25	0.90	6.44e+005	3.14	Base To Base		
5	I6-534006d\0070.wiff	252.41	0.96	6.44e+005	3.16	Base To Base		
6	I6-534006d\0071.wiff	244.92	-2.0	6.28e+005	3.16	Base To Base		
7	I6-534006d\0072.wiff	752.43	0.32	1.73e+006	3.14	Base To Base	1 = -	
8	l6-534006d\0073.wiff	751.32	0.18	1.73e+006	3.16	Base To Base	1 7	
9	I6-534006d\0074.wiff	747.66	-0.31	1.72e+006	3.13	Base To Base		
10	l6-534006d\0075.wiff	994.10	-0.59	2.24e+006	3.14	Base To Base		
11	I6-534006d\0076.wiff	1007.3	0.73	2.27e+006	3.12	Base To Base	<del>                                     </del>	
12	I6-534006d\0077.wiff	1012.6	1.3	2.28e+006	3.13	Base To Base		
13	I6-534006d\0078.wiff	1976.0	-1.2	4.16e+006	3.11	Base To Base	┪	
14	I6-534006d\0079.wiff	1971.1	-1.4	4.15e+006	3.14	Base To Base	1 5	······································
15	l6-534006d\0080.wiff	1991.8	-0.41	4.19e+006	3.15	Base To Base		
16	I6-534006d\0081.wiff	2999.1	-0.029	5.93e+006	3.13	Base To Base	一一	
17	I6-534006d\0082.wiff	2995.5	-0.15	5.93e+006	3.13	Base To Base	1-5	
18	I6-534006d\0083.wiff	3056.0	1.9	6.03e+006	3.10	Base To Base		······
19	l6-534006d\0084.wiff	No Peak	0.0	0.00e+000	0.00	No Peak		percent noise change
20	l6-534006d\0085.wiff	998.37	-0.16	2.25e+006	3.14	Base To Base		
21	I6-534006d\0086.wiff	1021.3	2.1	2.29e+006	3.09	Base To Base		***************************************
22	I6-534006d\0087.wiff	1017.2	1.7	2.29e+006	3.10	Base To Base		······································

Printing Date: Thursday, December 07, 2006 Printing Time: 9:17:59 AM

Operator: ullman Analyst Version: 1.4.1







Printing Date: Thursday, December 07, 2006 Printing Time: 9:19:29 AM Operator: ullman Analyst Version: 1.4.1 Table A-2: 16-534006e Data

Note: Validation Session II QC samples are listed as the diluted concentration of QC 1000.

	File Name	Sample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record	Analyte Concentration (ng/mL)	Dilution Factor	Calculated Concentration (ng/mL)
1	I6-534006e\0088.wiff	6-1	Diluent	Unknown		(X) (X) (X)	N/A	1.00	No Peak
2	I6-534006e\0089.wiff	11-2	C 250	Standard		$\boxtimes$	250.00	1.00	252.80
3	I6-534006e\0090.wiff	11-3	C 250	Standard		$\boxtimes$	250.00	1.00	247.04
4	I6-534006e\0091.wiff	11-4	C 250	Standard		$\boxtimes$	250.00	1.00	249.71
5	I6-534006e\0092.wiff	11-5	C 750	Standard		$\boxtimes$	750.00	1.00	757.33
3	I6-534006e\0093.wiff	11-6	C 750	Standard		X	750.00	1.00	739.40
7	I6-534006e\0094.wiff	11-7	C 750	Standard		$\boxtimes$	750.00	1.00	743.18
3	I6-534006e\0095.wiff	11-8	C 1000	Standard		$\boxtimes$	1000.0	1.00	1011.8
)	I6-534006e\0096.wiff	11-9	C 1000	Standard		$\boxtimes$	1000.0	1.00	1036.3
10	I6-534006e\0097.wiff	11-10	C 1000	Standard		$\boxtimes$	1000.0	1.00	996.03
11	l6-534006e\0098.wiff	11-11	C 2000	Standard		$\boxtimes$	2000.0	1.00	1911.9
12	I6-534006e\0099.wiff	11-12	C 2000	Standard	1	$\boxtimes$	2000.0	1.00	1936.3
13	I6-534006e\0100.wiff	11-13	C 2000	Standard		×	2000.0	1.00	2037.0
14	I6-534006e\0101.wiff	11-14	C 3000	Standard		$\boxtimes$	3000.0	1.00	2969.5
15	I6-534006e\0102.wiff	11-15	C 3000	Standard		⊠	3000.0	1.00	3051.5
16	I6-534006e\0103.wiff	11-16	C 3000	Standard		$\boxtimes$	3000.0	1.00	3068.2
7	I6-534006e\0104.wiff	12-6	QC 0	Quality Control		⊠	0.0000	1.00	No Peak
8	I6-534006e\0105.wiff	12-7	QC 1000	Quality Control		$\boxtimes$	1000.0	1.00	1117.6
9	I6-534006e\0106.wiff	12-8	QC 1000	Quality Control		×	1000.0	1.00	1141.2
20	I6-534006e\0107.wiff	12-9	QC 1000	Quality Control		×	1000.0	1.00	1134.7

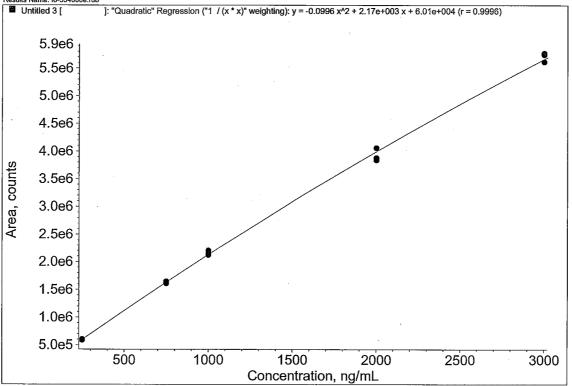
-31-73 of 281

	File Name	%RE	Analyte Peak Area (counts)	Analyte Retention Time (min)	Analyte Integration Type	Record Modified	Sample Annotation
1	I6-534006e\0088.wiff	N/A	0.00e+000	0.00	No Peak		percent noise change
2	I6-534006e\0089.wiff	1.1	6.01e+005	3.11	Base To Base		<u> </u>
3	I6-534006e\0090.wiff	-1.2	5.89e+005	3.12	Base To Base	1 7	
4	I6-534006e\0091.wiff	-0.11	5.95e+005	3.10	Base To Base		<del>                                     </del>
5	I6-534006e\0092.wiff	0.98	1.64e+006	3.10	Base To Base		<u> </u>
6	I6-534006e\0093.wiff	-1.4	1.61e+006	3.11	Base To Base		<u> </u>
7	l6-534006e\0094.wiff	-0.91	1.61e+006	3.11	Base To Base		
8	I6-534006e\0095.wiff	1.2	2.15e+006	3.12	Base To Base		<u> </u>
9	I6-534006e\0096.wiff	3.6	2.20e+006	3.10	Base To Base		
10	l6-534006e\0097.wiff	-0.40	2.12e+006	3.10	Base To Base		
11	I6-534006e\0098.wiff	-4.4	3.84e+006	3.11	Base To Base		
12	l6-534006e\0099.wiff	-3.2	3.88e+006	3.10	Base To Base		
13	I6-534006e\0100.wiff	1.8	4.06e+006	3.11	Base To Base		
14	I6-534006e\0101.wiff	-1.0	5.61e+006	3.10	Valley		
15	I6-534006e\0102.wiff	1.7	5.74e+006	3.08	Base To Base		
16	I6-534006e\0103.wiff	2.3	5.77e+006	3.11	Base To Base		
17	I6-534006e\0104.wiff	0.0	0.00e+000	0.00	No Peak		noise percent change
18	I6-534006e\0105.wiff	12.	2.36e+006	3.10	Base To Base		
19	I6-534006e\0106.wiff	14.	2.40e+006	3.09	Base To Base		
	I6-534006e\0107.wiff	13.	2.39e+006	3.10	Base To Base	Τп	

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Table A-3: 16-534006f Data

Note: Validation Session III QC samples are listed as the diluted concentration of QC 1000.

-34-

76 of 281

Results Path: D:\SciexData\Projects\534006\Dosing\Results\l6-534006f.rdb
Results Name: I6-534006f.rdb

Page 1 of 4

	File Name	Sample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record	Analyte Concentration (ng/mL)	Dilution Factor	Calculated Concentration (ng/mL)
1	I6-534006f\0108.wiff	15-2	system suit	Unknown	l-	20000000000	N/A	1.00	214.62
2	I6-534006f\0109.wiff	15-2	system suit	Unknown	***************************************		N/A	1.00	222.06
3	I6-534006f\0110.wiff	15-2	system suit	Unknown			N/A	1.00	239.86
4	I6-534006f\0111.wiff	na	mobile phase	Unknown	***************************************		N/A	1.00	No Peak
5	I6-534006f\0112.wiff	15-2	C 250	Standard /	***************************************		250.00	1.00	256.01
6	I6-534006f\0113.wiff	15-3	C 250	Standard	***************************************		250.00	1.00	241.89
7	I6-534006f\0114.wiff	15-4	C 250	Standard	•		250.00	1.00	252.08
8	I6-534006f\0115.wiff	15-5	C 750	Standard	***************************************	$\square$	750.00	1.00	749.77
9	I6-534006f\0116.wiff	15-6	C 750	Standard	***************************************	$\boxtimes$	750.00	1.00	730.09
10	I6-534006f\0117.wiff	15-7	C 750	Standard	•••••	$\boxtimes$	750.00	1.00	757.49
11	l6-534006f\0118.wiff	15-8	C 1000	Standard	***************************************		1000.0	1.00	997.99
12	I6-534006f\0119.wiff	15-9	C 1000	Standard			1000.0	1.00	1018.6
13	I6-534006f\0120.wiff	15-10	C 1000	Standard	terino anti-		1000.0	1.00	1017.2
14	I6-534006f\0121.wiff	15-11	C 2000	Standard		$\square$	2000.0	1.00	1952.9
15	I6-534006f\0122.wiff	15-12	C 2000	Standard	***************************************	Ø	2000.0	1.00	1962.0
16	16-534006f\0123.wiff	15-13	C 2000	Standard			2000.0	1.00	2020.1
17	I6-534006f\0124.wiff	15-14	C 3000	Standard	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,		3000.0	1.00	3007.7
18	I6-534006f\0125.wiff	15-15	C 3000	Standard			3000.0	1.00	3026.7
19	I6-534006f\0126.wiff	15-16	C 3000	Standard	***************************************		3000.0	1.00	3015.6
20	I6-534006f\0127.wiff	na	mobile phase	Unknown	***************************************		N/A	1.00	No Peak
21	I6-534006f\0128.wiff	16-6	QC 0	Quality Control	***************************************	$\square$	0.0000	1.00	No Peak
22	I6-534006f\0129.wiff	16-7	QC 1000	Quality Control	**********	$\square$	1000.0	1.00	1015.4
23	I6-534006f\0130.wiff	16-8	QC 1000	Quality Control			1000.0	1.00	1019.4
24	I6-534006f\0131.wiff	16-9	QC 1000	Quality Control	***************************************		1000.0	1.00	1010.0
25	I6-534006f\0132.wiff	na	mobile phase	Unknown	***************************************	T	N/A	1.00	No Peak

Printing Date: Friday, December 08, 2006 Printing Time: 9:12:49 AM

Results Path: D:\SciexData\Projects\534006\Dosing\Results\16-534006f.rdb Results Name: 16-534006f.rdb

Page 2 of 4

	File Name	Sample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record	Analyte Concentration (ng/mL)	Dilution Factor	Calculated Concentration (ng/mL)
26	I6-534006f\0133.wiff	17-5	group	Unknown		1085,453,688	N/A	1.00	1220.1
27	I6-534006f\0134.wiff	17-6	group	Unknown	***************************************	T	N/A	1.00	1032.7
28	I6-534006∩0135.wiff	17-7	group	Unknown	•	1	N/A	1.00	1045.7
29	I6-534006N0136.wiff	17-8	group	Unknown			N/A	1.00	1060.5
30	I6-534006f\0137.wiff	na	mobile phase	Unknown	<del></del>		N/A	1.00	No Peak
31	I6-534006f\0138.wiff	8-2	C 250 PSS	Unknown		İ	N/A	1.00	259.95
32	16-534006↑0139.wiff	8-3	C 250 PSS	Unknown	***************************************	1	N/A	1.00	263.09
33	l6-534006f\0140.wiff	8-4	C 250 PSS	Unknown	*********		N/A	1.00	258.95
34	I6-534006f\0141.wiff	9-7	QC 1000 PSS	Unknown	***************************************	1	N/A	1.00	1019.2
35	I6-534006f\0142.wiff	9-8	QC 1000 PSS	Unknown		1	N/A	1.00	1040.2
36	I6-534006f\0143.wiff	9-9	QC 1000 PSS	Unknown	Anabranens	1	N/A	1.00	1034.7
37	I6-534006f\0144.wiff	8-14	C 3000 PSS	Unknown	********		N/A	1.00	3176.8
38	I6-534006f\0145.wiff	8-15	C 3000 PSS	Unknown	***************************************	1	N/A	1.00	3137.0
39	I6-534006f\0146.wiff	8-16	C 3000 PSS	Unknown			N/A	1.00	3217.3
40	I6-534006f\0147.wiff	na	mobile phase	Unknown	***************************************	1	N/A	1.00	No Peak

Printing Date: Friday, December 08, 2006 Printing Time: 9:12:49 AM

	File Name	%RE	Analyte Peak Area (counts)	Analyte Retention Time (min)	Analyte Integration Type	Record Modified	Sample Annotation
1	I6-534006f\0108.wiff	N/A	6.12e+005	3.39	Base To Base		RT & peak split change
2	I6-534006f\0109.wiff	N/A	6.30e+005	3.30	Base To Base		
3	I6-534006f\0110.wiff	N/A	6.74e+005	3.28	Base To Base		
4	I6-534006f\0111.wiff	N/A	0.00e+000	0.00	No Peak	$\boxtimes$	percent noise
5	I6-534006f\0112.wiff	2.4	7.13e+005	3.30	Base To Base		
6	I6-534006f\0113.wiff	-3.2	6.79e+005	3.27	Base To Base		
7	l6-534006f\0114.wiff	0.83	7.03e+005	3.27	Base To Base		
8	l6-534006f\0115.wiff	-0.031	1.87e+006	3.24	Base To Base		
9	I6-534006f\0116.wiff	-2.7	1.82e+006	3.24	Base To Base		***************************************
10	l6-534006f\0117.wiff	1.0	1.89e+006	3.24	Base To Base		
11	l6-534006f\0118.wiff	-0.20	2.42e+006	3.26	Base To Base		
12	I6-534006f\0119.wiff	1.9	2.46e+006	3.24	Base To Base		
13	I6-534006f\0120.wiff	1.7	2.46e+006	3.23	Base To Base		
14	I6-534006f\0121.wiff	-2.4	4.35e+006	3.23	Base To Base		
15	16-534006f\0122.wiff	-1.9	4.37e+006	3.21	Base To Base		
16	I6-534006f\0123.wiff	1.0	4.48e+006	3.22	Base To Base		peak split
17	16-534006f\0124.wiff	0.26	6.15e+006	3.22	Base To Base		***************************************
18	I6-534006f\0125.wiff	0.89	6.18e+006	3.21	Base To Base		
19	I6-534006f\0126.wiff	0.52	6.16e+006	3.22	Base To Base		
20	I6-534006f\0127.wiff	N/A	0.00e+000	0.00	No Peak		percent noise
21	l6-534006f\0128.wiff	0.0	0.00e+000	0.00	No Peak		percent noise
22	I6-534006f\0129.wiff	1.5	2.46e+006	3.23	Base To Base		
23	I6-534006f\0130.wiff	1.9	2.47e+006	3.22	Base To Base		
24	I6-534006f\0131.wiff	1.0	2.45e+006	3.20	Base To Base		
25	I6-534006f\0132.wiff	N/A	0.00e+000	0.00	No Peak		percent noise

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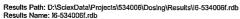
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-37

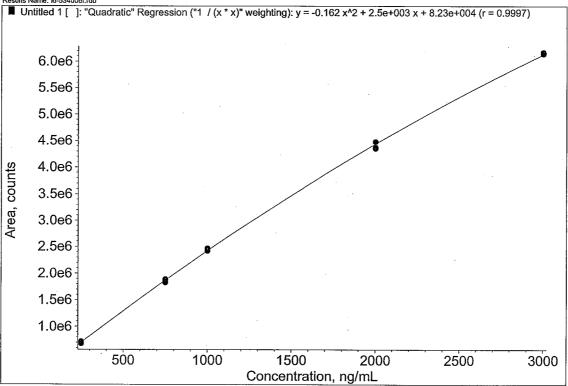
79 of 281

	File Name	%RE	Analyte Peak Area (counts)	Analyte Retention Time (min)	Analyte Integration Type	Record Modified	Sample Annotation
26	I6-534006f\0133.wiff	N/A	2.90e+006	3.22	Base To Base	+	
27	I6-534006f\0134.wiff	N/A	2.50e+006	3.21	Base To Base		
28	I6-534006f\0135.wiff	N/A	2.52e+006	3.21	Base To Base	十一一	
29	I6-534006f\0136.wiff	N/A	2.56e+006	3.18	Base To Base	1 7	<u> </u>
30	I6-534006f\0137.wiff	N/A	0.00e+000	0.00	No Peak		percent noise
31	I6-534006f\0138.wiff	N/A	7.22e+005	3.17	Base To Base	1 <u>5</u> -	
32	l6-534006f\0139.wiff	N/A	7.30e+005	3.22	Base To Base		
33	l6-534006⋀0140.wiff	N/A	7.20e+005	3.17	Base To Base	1 7	
34	I6-534006N0141.wiff	N/A	2.47e+006	3.20	Valley		
35	I6-534006f\0142.wiff	N/A	2.51e+006	3.21	Base To Base		
36	I6-534006f\0143.wiff	N/A	2.50e+006	3.18	Base To Base	1 7	
37	I6-534006f\0144.wiff	N/A	6.40e+006	3.16	Base To Base	1 -	
38	16-534006f\0145.wiff	N/A	6.34e+006	3.18	Valley	$\Box$	
39	I6-534006f\0146.wiff	N/A	6.46e+006	3.19	Base To Base	$\vdash \overline{\sqcap}$	
40	I6-534006f\0147.wiff	N/A	0.00e+000	0.00	No Peak		percent noise

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Page 1 of 1



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Table A-4: I6-534006g Data

Note:
Validation Session IV
QC samples are listed as the diluted concentration of QC 1000.

-40-

82 of 281

	File Name	Sample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record	Analyte Concentration (ng/mL)	Dilution Factor
	16-534006g\0148.wiff	22-2	system suit	Unknown	t		N/A	1.00
2	l6-534006g\0149.wiff	22-2	system suit	Unknown			N/A	1.00
3	l6-534006g\0150.wiff	22-2	system suit	Unknown	***************************************		N/A	1.00
	l6-534006g\0151.wiff	na	mobile phase	Unknown			N/A	1.00
	16-534006g\0152.wiff	22-2	C 250	Standard	***************************************	$\boxtimes$	250.00	1.00
332	l6-534006g\0153.wiff	22-3	C 250	Standard	ī ·		250.00	1.00
<b>1</b> 2-, 0	l6-534006g\0154.wiff	22-4	C 250	Standard		×	250.00	1.00
3	16-534006g\0155.wiff	22-5	C 750	Standard		$\boxtimes$	750.00	1.00
)	l6-534006g\0156.wiff	22-6	C 750	Standard		$\boxtimes$	750.00	1.00
0	l6-534006g\0157.wiff	22-7	C 750	Standard	1	図	750.00	1.00
1	l6-534006g\0158.wiff	22-8	C 1000	Standard	1	$\boxtimes$	1000.0	1.00
2	16-534006g\0159.wiff	22-9	C 1000	Standard	ji -	$\boxtimes$	1000.0	1.00
3	l6-534006g\0160.wiff	22-10	C 1000	Standard	Ţi —	×	1000.0	1.00
4	l6-534006g\0161.wiff	22-11	C 2000	Standard	† —	×	2000.0	1.00
5	l6-534006g\0162.wiff	22-12	C 2000	Standard	i	×	2000.0	1.00
6	16-534006g\0163.wiff	22-13	C 2000	Standard	1	×	2000.0	1.00
7	l6-534006g\0164.wiff	22-14	C 3000	Standard	•	×	3000.0	1.00
8	16-534006g\0165.wiff	22-15	C 3000	Standard	<u></u>	$\boxtimes$	3000.0	1.00
9	16-534006g\0166.wiff	22-16	C 3000	Standard	***************************************	$\boxtimes$	3000.0	1.00
0	l6-534006g\0167.wiff	na	mobile phase	Unknown	ļ		N/A	1.00
105	l6-534006g\0168.wiff	23-6	QC 0	Quality Control	***************************************	Ø	0.0000	1.00
2	l6-534006g\0169.wiff	23-7	QC 1000	Quality Control	• • • • • • • • • • • • • • • • • • • •		1000.0	1.00
	l6-534006g\0170.wiff	23-8	QC 1000	Quality Control	***************************************	$\boxtimes$	1000.0	1.00
24	l6-534006g\0171.wiff	23-9	QC 1000	Quality Control			1000.0	1.00
25	l6-534006g\0172.wiff	na	mobile phase	Unknown	*		N/A	1.00

Printing Date: Tuesday, December 12, 2006 Printing Time: 12:57:08 PM

	File Name	Sample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record	Analyte Concentration (ng/mL)	Dilution Factor	
26	I6-534006g\0173.wiff	24-3	group	Unknown			N/A	1.00	
7	I6-534006g\0174.wiff	24-4	group	Unknown			N/A	1.00	
8:	l6-534006g\0175.wiff	na	mobile phase	Unknown			N/A	1.00	
								٠	

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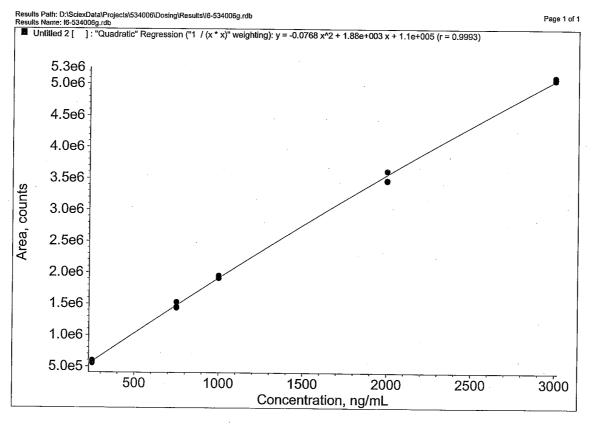
							manuscramina de la companya della companya della companya de la companya della co	
	File Name	Calculated Concentration (ng/mL)	%RE	Analyte Peak Area (counts)	Analyte Retention Time (min)	Analyte Integration Type	Record	Sample Annotation
-	l6-534006g\0148.wiff	240.60	N/A	5.58e+005	4.08	Base To Base		
2	I6-534006g\0149.wiff	271.92	N/A	6.15e+005	4.08	Base To Base		
3	I6-534006g\0150.wiff	260.12	N/A	5.94e+005	4.03	Base To Base		
4	l6-534006g\0151.wiff	No Peak	N/A	0.00e+000	0.00	No Peak		***************************************
2	l6-534006g\0152.wiff	259.47	3.8	5.92e+005	4.11	Base To Base		***************************************
9	l6-534006g\0153.wiff	251.90	0.76		4.10	Base To Base		
7	I6-534006g\0154.wiff	238.33	4.7	5.53e+005	4.10	Valley		***************************************
8	I6-534006g\0155.wiff	726.42	-3.1	1.43e+006	4.10	Base To Base		
6	I6-534006g\0156.wiff	737.07	-1.7	1.45e+006	4.08	Base To Base		***************************************
10	l6-534006g\0157.wiff	778.23	3.8	1.53e+006	4.08	Base To Base		***************************************
1	I6-534006g\0158.wiff	997.70	-0.23	1.91e+006	4.10	Base To Base		***************************************
12	I6-534006g\0159.wiff	1022.0	2.2	1.95e+006	4.09	Base To Base		was a second of the second of
<del>.</del>	I6-534006g\0160.wiff	1014.4	1.4	1.94e+006	4.05	Base To Base		
14	l6-534006g\0161.wiff	1937.1	-3.1	3.46e+006	4.08	Base To Base		
15	I6-534006g\0162.wiff	1943.0	-2.9	3.47e+006	4.09	Base To Base		
16	I6-534006g\0163.wiff	2034.9	1.7	3.62e+006	4.08	Base To Base		***************************************
17	I6-534006g\0164.wiff	3039.3	1.3	5.11e+006	4.04	Base To Base		
18	I6-534006g\0165.wiff	3005.5	0.18	5.06e+006	4.10	Base To Base		***************************************
19	16-534006g\0166.wiff	3019.2	0.64	5.08e+006	4.07	Base To Base		arteretis et defenses desse stress desse de la cons
20	l6-534006g\0167.wiff	No Peak	N/A	0.00e+000	0.00	No Peak		
21	I6-534006g\0168.wiff	No Peak	0.0	0.00e+000	0.00	No Peak		
72	I6-534006g\0169.wiff	1002.3	0.23	1.92e+006	4.03	Base To Base		
23	I6-534006g\0170.wiff	1019.7	2.0	1.95e+006	4.08	Base To Base		
24	I6-534006g\0171.wiff	1003.5	0.35	1.92e+006	4.05	Base To Base		
22	l6-534006g\0172.wiff	No Peak	N/A	0.00e+000	0.00	No Peak		

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	File Name	Calculated Concentration (ng/mL)	%RE	Analyte Peak Area (counts)	Analyte Retention Time (min)	Analyte Integration Type	Record Modified	Sample Annotation
6	I6-534006g\0173.wiff	1081.3	N/A	2.05e+006	4.05	Valley		
7	I6-534006g\0174.wiff	1063.8	N/A	2.02e+006	4.09	Base To Base		***************************************
В	l6-534006g\0175.wiff	No Peak	N/A	0.00e+000	0.00	No Peak		,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,
							•	
	•							
								•
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# APPENDIX C

Pretest Clinical Observations

PROJECT NO.:534006P		COKINETIC AND EXCRETION STUDY IN RATS CAL FINDINGS: TOTAL OCCURRENCE/NO. OF ANIMALS	PAGE 1
		MALE	
	TABLE RANGE: GROUP:	12-01-06 TO 12-11-06	1
NORMAL -NO SIGNIFICANT CLINICAL	OBSERVATIONS		29/15
EXCRETA -SOFT FECES			1/ 1
1- PRETEST			

PROJECT NO.:534006P

# A PHARMACOKINETIC AND EXCRETION STUDY IN RATS SUMMARY OF CLINICAL FINDINGS: TOTAL OCCURRENCE/NO. OF ANIMALS

PAGE

	F E M A L E	
TABLE RANGE: GROUP:	12-01-06 TO 12-11-06	1
NORMAL -NO SIGNIFICANT CLINICAL OBSERVATIONS		30/15
1- PRETEST		

PCSUv4.07 06/18/2007

# APPENDIX D

Animal Room Environmental Conditions

PROJECT NO.

534006

#### A PHARMACOKINETIC AND EXCRETION STUDY IN RATS TEMPERATURE/HUMIDITY - DAILY SUMMARY REPORT BY STUDY

STUDY SPECIFICATIONS: 534006 DATE IN: 11/28/06 TIME IN: 7:00 DATE OUT: 12/13/06 TIME OUT: 16:00

ROOM SPECIFICATIONS: B ROOM 127 LOW TEMPERATURE °F: 66.0 HIGH TEMPERATURE °F: 76.0 LOW HUMIDITY: 30.0 SPECIES: RAT LOW TEMPERATURE °C: 18.9 HIGH TEMPERATURE °C: 24.4 HIGH HUMIDITY: 70 0

	1412		HOW TENEDIN	MIUNE C.	10.9 HIGH TEMPERATURE	C: 24.4 HIGH HUMIDITY: /(	J. U
	TEMP	ERATURE	HUMI	DITY			
DATE	MEAN (°F)	MEAN (°C)	MEAN (9	%RH)			
28-Nov-06	70.6	21.4	33.6				
29-Nov-06	70.5	21.4	45.9				
30-Nov-06	70.5	21.4	44.5				
01-Dec-06	70.4	21.3	31.6				
02-Dec-06	70.5	21.4	29.3 -	•	•		
03-Dec-06	70.5	21.4	37.2				
04-Dec-06	70.4	21.4	36.2				
05-Dec-06	70.5	21.4	37.4			•	
06-Dec-06	70.5	21.4	39.4				
07-Dec-06	70.5	21.4	35.2				
08-Dec-06	70.2	21.2	36.4				
09-Dec-06	70.5	21.4	34.3				
10-Dec-06	70.5	21.4	36.9	•			
11-Dec-06	70.6	21.4	38.2				
12-Dec-06	70.4	21.4	41.6				
13-Dec-06	70.5	21.4	43.2				

NOTE: + = VALUE WAS GREATER THAN HIGH RANGE

NOTE: MEANS REPRESENT THE MEAN OF THE DAILY VALUES

VERSION 1.09 2/22/2007 12:44 PAGE 1

92 of 281

- = VALUE WAS LESS THAN LOW RANGE

REPORT 4

PROJECT NO.

534006

#### A PHARMACOKINETIC AND EXCRETION STUDY IN RATS TEMPERATURE/HUMIDITY - DAILY SUMMARY REPORT BY STUDY

STUDY SPECIFICATIONS:

DATE OUT: 12/13/06

7:00 16:00

ROOM SPECIFICATIONS:

B ROOM 127

534006

LOW TEMPERATURE °F: 66.0 HIGH TEMPERATURE °F: 76.0

LOW HUMIDITY: 30.0

SPECIES:

RAT

LOW TEMPERATURE °C: 18.9 HIGH TEMPERATURE °C: 24.4

PAGE 2

HIGH HUMIDITY: 70.0

TEMPERATURE

HUMIDITY

DATE

MEAN (°F) MEAN (°C) MEAN (%RH)

GRAND STATS	MEAN	MIN	MAX
TEMPERATURE °F	70.5	70.2	70.6
TEMPERATURE °C	21.4	21.2	21.4
HUMIDITY (%RH)	37.6	29.3 -	45.9
N DAVS	16		

93 of 281

NOTE: + = VALUE WAS GREATER THAN HIGH RANGE

- = VALUE WAS LESS THAN LOW RANGE

NOTE: MEANS REPRESENT THE MEAN OF THE DAILY VALUES

REPORT 4 VERSION 1.09

2/22/2007 12:44

PROJECT NO.

534006

A PHARMACOKINETIC AND EXCRETION STUDY IN RATS TEMPERATURE/HUMIDITY - END OF STUDY SUMMARY REPORT

12:48 22-Feb-07

PAGE 1

ROOM SPECIFICATIONS: B ROOM 127

SPECIES:

94 of 281

RAT

LOW TEMPERATURE: 66.0 76.0

11/28/06 DATE IN:

HIGH TEMPERATURE: LOW HUMIDITY:

TIME IN: 7:00

HIGH HUMIDITY:

DATE OUT: 12/13/06 16:00

30.0 70.0 TIME OUT:

TEMPERATURE

HUMIDITY

37.5

14.8

54.2

6.15

370

ROOM B ROOM 127 SUMMARY

MEAN MIN

MAX

SD

70.5 64.7

73.9 0.61

370 N SAMPLES FIRST DAY 11/28/06

LAST DAY 12/13/06 N DAYS 16

NOTE: TEMPERATURE UNITS = DEGREES FAHRENHEIT

HUMIDITY UNITS = % RELATIVE HUMIDITY

NOTE: MEANS REPRESENT THE MEAN OF ALL VALUES

REPORT 5 VERSION 1.10

2/22/2007 12:48

A PHARMACOKINETIC AND EXCRETION STUDY IN RATS TEMPERATURE/HUMIDITY - END OF STUDY SUMMARY REPORT

PROJECT NO.

534006

12:48 22-Feb-07

PAGE 2

STUDY 534006 SUMMARY

MEAN	70.5	37.5
MIN	64.7	14.8
MAX	73.9	54.2
SD	0.61	6.15
N SAMPLES	370	370
FIRST DAY	11/28/06	
LAST DAY	12/13/06	
N DAYS	16	

95 of 281

NOTE: TEMPERATURE UNITS = DEGREES FAHRENHEIT HUMIDITY UNITS = % RELATIVE HUMIDITY

NOTE: MEANS REPRESENT THE MEAN OF ALL VALUES

REPORT 5 VERSION 1.10 2/22/2007 12:48

## APPENDIX E

Bioanalytical Report [

# PHARMACOKINETIC (IN BLOOD) AND EXCRETION STUDY OF [ ] IN RATS

Analyses Of Biological Samples

-1-

]

# TABLE OF CONTENTS

	<u>Page</u>
	Table Of Contents2
	Index Of Tables3
	Index Of Figures4
	Index Of Attachments8
1.	Introduction9
2.	Experimental10
2.1.	Identification Of Blank Serum And Urine
2.2.	Instruments
2.2.1.	High Performance Liquid Chromatography
2.3.	Mass Spectrometry (API 4000)
2.4.	Mass Spectrometry (API 5000)12
2.5.	Preparation Of Mobile Phase14
2.6.	Preparation Of The [ ] Calibration Stock Solution - 0.1 mg/mL14
2.7.	Preparation Of The [ ] Calibration Stock Solution - 1.0 mg/mL14
2.8.	Preparation Of The [ ] Quality Control Stock Solution14
2.9.	Preparation Of Calibration Samples
2.10.	Preparation Of Quality Control Samples
2.11.	Sample Processing
2.12.	Concentration Quantitation 16
3.	Results And Discussion
3.1.	Method Validation
3.2.	Sensitivity
3.3.	Specificity/Selectivity
3.4.	Ruggedness 17
3.5.	Calibration Acceptability
3.6.	Accuracy And Precision
3.7.	Stability

	<u>Page</u>					
3.7.1.	Stability Of [ ] In Refrigerated Stock Solutions					
3.7.2.	Long Term Frozen Stability Of [ ] In Serum And Urine Samples20					
3.7.3.	Stability Of [ ] In Serum And Urine Samples At Room Temperature					
3.7.4.	Freeze-Thaw Stability Of [ ] In Serum And Urine Samples21					
3.8.	Analysis Of Experimental Samples					
4.	Conclusion					
5.	Key Study Personnel And Report Submission23					
	INDEX OF TABLES					
Table 1:	[ ] Back-Calculated Concentrations And Intra- And Inter-Session Statistics Of Calibration Samples In Rat Serum25					
Table 2:	[ ] Back-Calculated Concentrations And Intra-Session Statistics Of Calibration Samples In Rat Urine					
Table 3:	[ ] Concentrations And Intra- And Inter-Session Statistics Of Quality Control Samples In Rat Serum					
Table 4:	[ ] Concentrations And Intra-Session Statistics Of Quality Control Samples In Rat Urine					
Table 5:	[ ] Concentrations And Intra-Session Statistics Of Quality Control Samples In Cage Rinse					
Table 6:	Refrigerated (4°C) Stability Of [ ] In DI Water Stock Solutions34					
Table 7:	Stability Of [ ] In Rat Serum - Long-Term Frozen (-20°C) Storage					
Table 8:	Stability Of [ ] In Rat Urine - Long-Term Frozen (-20°C) Storage					
Table 9:	Room Temperature Stability Of [ ] In Rat Serum					
Table 10:	Room Temperature Stability Of [ ] In Rat Urine					

		Page
Table 11:	Freeze-Thaw Stability Of [ ] In Rat Serum	39
Table 12:	Freeze-Thaw Stability Of [ ] In Rat Urine	40
Table 13:	Rat Serum Experimental Sample [ ] Concentration Data	42
Table 14:	Rat Urine/Cage Rinse Experimental Sample [ ] Concentration Data	44
Table 15:	[ ] Calibration Samples - Routine Analyses	45
Table 16:	[ ] Quality Control Samples - Routine Analyses	52
	INDEX OF FIGURES	
Figure 1:	Representative Chromatogram Of A Solvent Blank	57
Figure 2:	Representative Chromatogram Of A Processed Blank Rat Serum Sample	57
Figure 3:	Representative Chromatogram Of A Processed 10.0 ng/mL Rat Serum Calibration Sample	58
Figure 4:	Representative Chromatogram Of A Processed 30.0 ng/mL Rat Serum Calibration Sample	58
Figure 5:	Representative Chromatogram Of A Processed 100 ng/mL Rat Serum Calibration Sample	59
Figure 6:	Representative Chromatogram Of A Processed 300 ng/mL Rat Serum Calibration Sample	59
Figure 7:	Representative Chromatogram Of A Processed 500 ng/mL Rat Serum Calibration Sample	60
Figure 8:	Representative Chromatogram Of A Processed 750 ng/mL Rat Serum Calibration Sample	60

		<u>Page</u>
Figure 9:	Representative Chromatogram Of A Processed 1000 ng/mL Rat Serum Calibration Sample	61
Figure 10:	Representative Chromatogram Of A Processed 30.0 ng/mL Rat Serum QC Sample	61
Figure 11:	Representative Chromatogram Of A Processed 250 ng/mL Rat Serum QC Sample	62
Figure 12:	Representative Chromatogram Of A Processed 750 ng/mL Rat Serum QC Sample	62
Figure 13:	Representative Chromatogram Of A Processed 30,000 ng/mL Rat Serum Dilutional QC Sample	63
Figure 14:	Chromatogram Of A Processed Group 1 Male Rat Serum Sample On Study Day 0, Pre-Dose	63
Figure 15:	Chromatogram Of A Processed Group 1 Female Rat Serum Sample On Study Day 0, Pre-Dose	64
Figure 16:	Chromatogram Of A Processed Group 1 Male Rat Serum Sample On Study Day 0, 2 Minutes Post-Dose	64
Figure 17:	Chromatogram Of A Processed Group 1 Female Rat Serum Sample On Study Day 0, 2 Minutes Post-Dose	65
Figure 18:	Chromatogram Of A Processed Group 1 Male Rat Serum Sample On Study Day 0, 10 Minutes Post-Dose	65
Figure 19:	Chromatogram Of A Processed Group 1 Female Rat Serum Sample On Study Day 0, 10 Minutes Post-Dose	66
Figure 20:	Chromatogram Of A Processed Group 1 Male Rat Serum Sample On Study Day 0, 20 Minutes Post-Dose	66
Figure 21:	Chromatogram Of A Processed Group 1 Female Rat Serum Sample On Study Day 0, 20 Minutes Post-Dose	67
Figure 22:	Chromatogram Of A Processed Group 1 Male Rat Serum Sample On Study Day 0, 30 Minutes Post-Dose	67

		<u>Page</u>
Figure 23:	Chromatogram Of A Processed Group 1 Female Rat Serum Sample On Study Day 0, 30 Minutes Post-Dose	68
Figure 24:	Chromatogram Of A Processed Group 1 Male Rat Serum Sample On Study Day 0, 1 Hour Post-Dose	68
Figure 25:	Chromatogram Of A Processed Group 1 Female Rat Serum Sample On Study Day 0, 1 Hour Post-Dose	69
Figure 26:	Chromatogram Of A Processed Group 1 Male Rat Serum Sample On Study Day 0, 3 Hours Post-Dose	69
Figure 27:	Chromatogram Of A Processed Group 1 Female Rat Serum Sample On Study Day 0, 3 Hours Post-Dose	70
Figure 28:	Chromatogram Of A Processed Group 1 Male Rat Serum Sample On Study Day 0, 5 Hours Post-Dose	70
Figure 29:	Chromatogram Of A Processed Group 1 Female Rat Serum Sample On Study Day 0, 5 Hours Post-Dose	71
Figure 30:	Chromatogram Of A Processed Group 1 Male Rat Serum Sample On Study Day 0, 7 Hours Post-Dose	71
Figure 31:	Chromatogram Of A Processed Group 1 Female Rat Serum Sample On Study Day 0, 7 Hours Post-Dose	72
Figure 32:	Chromatogram Of A Processed Group 1 Male Rat Serum Sample On Study Day 0, 24 Hours Post-Dose	72
Figure 33:	Chromatogram Of A Processed Group 1 Female Rat Serum Sample On Study Day 0, 24 Hours Post-Dose	73
Figure 34:	Chromatogram Of A Processed Group 1 Male Rat Serum Sample On Study Day 0, 48 Hours Post-Dose	73
Figure 35:	Chromatogram Of A Processed Group 1 Female Rat Serum Sample On Study Day 0, 48 Hours Post-Dose	74
Figure 36:	Representative Chromatogram Of A Processed Blank Rat Urine Sample	74

		<u>Page</u>
Figure 37:	Representative Chromatogram Of A Processed 10.0 ng/mL Rat Urine Calibration Sample	75
Figure 38:	Representative Chromatogram Of A Processed 30.0 ng/mL Rat Urine Calibration Sample	75
Figure 39:	Representative Chromatogram Of A Processed 100 ng/mL Rat Urine Calibration Sample	76
Figure 40:	Representative Chromatogram Of A Processed 300 ng/mL Rat Urine Calibration Sample	76
Figure 41:	Representative Chromatogram Of A Processed 500 ng/mL Rat Urine Calibration Sample	77
Figure 42:	Representative Chromatogram Of A Processed 750 ng/mL Rat Urine Calibration Sample	77
Figure 43:	Representative Chromatogram Of A Processed 1000 ng/mL Rat Urine Calibration Sample	78
Figure 44:	Representative Chromatogram Of A Processed 30.0 ng/mL Rat Urine QC Sample	78
Figure 45:	Representative Chromatogram Of A Processed 250 ng/mL Rat Urine QC Sample	79
Figure 46:	Representative Chromatogram Of A Processed 750 ng/mL Rat Urine QC Sample	79
Figure 47:	Representative Chromatogram Of A Processed 30,000 ng/mL Rat Urine Dilutional QC Sample	80
Figure 48:	Representative Chromatogram Of A Processed 30.0 ng/mL Rat Cage Rinse QC Sample	80
Figure 49:	Representative Chromatogram Of A Processed 250 ng/mL Rat Cage Rinse QC Sample	81
Figure 50:	Representative Chromatogram Of A Processed 750 ng/mL Rat Cage Rinse QC Sample	81

		<u>Page</u>
Figure 51:	Representative Chromatogram Of A Processed 30,000 ng/mL Rat Cage Rinse Dilutional QC Sample	82
Figure 52:	Chromatogram Of A Processed Group 1 Rat Urine Sample On Study Day 0, 0-6 Hours Post-Dose	82
Figure 53:	Chromatogram Of A Processed Group 1 Rat Cage Rinse Sample On Study Day 0, 0-6 Hours Post-Dose	83
Figure 54:	Chromatogram Of A Processed Group 1 Rat Urine Sample On Study Day 0, 6-12 Hours Post-Dose	83
Figure 55:	Chromatogram Of A Processed Group 1 Rat Cage Rinse Sample On Study Day 0, 6-12 Hours Post-Dose	84
Figure 56:	Chromatogram Of A Processed Group 1 Rat Urine Sample On Study Day 0, 12-24 Hours Post-Dose	84
Figure 57:	Chromatogram Of A Processed Group 1 Rat Cage Rinse Sample On Study Day 0, 12-24 Hours Post-Dose	85
	INDEX OF ATTACHMENTS	
I:	Supporting Data	86

## 1. Introduction

This report provides a detailed description and validation of a high performance liquid chromatography tandem mass spectrometry (HPLC/MS/MS) method in the negative electrospray ionization (ESI-) mode for the determination 1 concentration in rat The method was validated over the concentration range of 10.0 to 1000 ng [ ]/mL of rat serum using a 50-µL sample. The method was cross-validated over the concentration range of 10.0 to 1000 ng [ /mL of rat urine using a 50-uL sample. Analyte stability in stock solutions and in stored rat serum and urine samples was assessed. [ ] in stock solutions was stable for at least 11 days of refrigerated storage. ] in rat serum was stable through at least 3 freeze-thaw cycles, for at least 4 hours of room temperature storage and for at least 124 days of frozen (approximately -20°C) I in rat urine was stable through at least 6 freeze-thaw cycles, for at least storage. 4 hours of room temperature storage and for at least 127 days of frozen (approximately -20°C) storage.

This report also details the analytical results for the determination of [ ] concentration in rat serum and urine samples. Analysis of experimental serum samples resulted in levels ranging from less than the lower limit of quantitation (<LLOQ; 10.0 ng/mL) to 100,720 ng [ ]/mL. Analysis of experimental urine samples resulted in total [ ] values ranging from 4.44 to 898  $\mu$ g. Analysis of experimental cage rinse samples resulted in total [ ] values ranging from 1.87 to 1006  $\mu$ g. The results of the sample analyses (Tables 13 and 14) and the supporting data (Attachment I) are included in this report.

The method for the determination of [ ] in rat serum, urine and cage rinse used acetonitrile (ACN) to de-proteinize 50  $\mu$ L of serum or urine. Following centrifugation, a portion of each supernatant fraction was collected, evaporated and reconstituted in mobile phase. The samples were analyzed with an HPLC/MS/MS assay using an ACE C8

column (50  $\times$  2.1 mm). The [ ] peak areas (y) and the theoretical concentrations of calibration samples were fit to a linear regression with  $1/x^2$  weighting, excluding the origin.

## 2. EXPERIMENTAL

## 2.1. IDENTIFICATION OF BLANK SERUM AND URINE

Blank rat serum and urine was purchased from Bioreclamation, East Meadow, New York. Blank serum and urine was stored frozen at approximately -20°C. Blank cage rinse was collected at [ ].

## 2.2. Instruments

The mass spectrometers used were an Applied Biosystems/MDS Sciex API 4000™ or an Applied Biosystems/MDS Sciex API 5000™ tandem quadrupole mass spectrometer (hereafter referred to as API 4000 or API 5000) equipped with a TurboIonSpray™ probe for ESI+ ionization. The HPLC systems used were an Agilent 1200 liquid chromatograph (used with the API 4000) equipped with an autosampler or a Spark Holland, Inc. Symbiosis Pharma System (used with the API 5000) equipped with an autosampler. Data acquisition and analysis were performed using Analyst® software version 1.4.2. The retention time, run time and mass spectrometer settings may have varied depending on column and mass spectrometer performance.

## 2.2.1. HIGH PERFORMANCE LIQUID CHROMATOGRAPHY

Analytical Column:

ACE C8,  $2.1 \times 50 \text{ mm}$ 

Column Temperature:

Ambient

20°C on Agilent 1200

Mobile Phase:

2 mM ammonium acetate in 70:30 (v/v)

deionized water: ACN

Detector:

Mass spectrometer with conditions as described in

Sections 2.3, and 2.4.

Flow Rate:

0.3 mL/minute

Injection Volume:

2 to 25 μL

-10-

106 of 281

Retention Time:

Approximately 3.0 to 3.5 minutes for [ ]

(Agilent 1200)

Approximately 4.6 to 6.2 minutes for [ ]

(Symbiosis)

Run Time:

Approximately 7.0 minutes (Agilent 1200)

Approximately 8.0 minutes (Symbiosis)

## 2.3. Mass Spectrometry (API 4000)

#### **Acquisition Parameters**

Scan Type:

Multiple reaction monitoring

Polarity:

Negative

Scan Mode:

Not Applicable

Ion Source:

Turbo Spray

Resolution Q1:

Unit

Resolution Q3:

Unit

Intensity Threshold:

0.00 cps

Settling Time:

0.00 msec

MR Pause:

5.0070 msec

MCA:

No

Step Size:

0.00 amu

Analyte	Q1 Mass (amu)	<u>Q3 Mass</u> (amu)	<u>Dwell</u> (msec)	<u>Parameter</u>	Start/Stop
[					]

DP = Declustering Potential

CE = Collision Energy

CXP = Collision Cell Exit Potential

#### 534006

#### Parameter Table

CUR (Curtain Gas): 40.00

GS1 (Gas 1): 60.00

GS2 (Gas 2): 60.00

IS (IonSpray Voltage): -4000.00

TEM (Temperature): 450.00

ihe (Interface Heater): OFF

CAD (Collisionally Activated 4.00 (nitrogen)

Dissociation Gas):

EP (Entrance Potential): -8.00

#### **Detector Parameters (Positive)**

CEM (Channel Electron 2000.0

Multiplier):

DF (Deflector): 300.0

# 2.4. MASS SPECTROMETRY (API 5000)

#### **Acquisition Parameters**

Scan Type: Multiple reaction monitoring

Polarity: Negative

Scan Mode: Not Applicable

Ion Source: Turbo Spray

Resolution Q1: Unit

Resolution Q1.

Resolution Q3: Unit

Intensity Threshold: 0.00 cps

Settling Time: 0.00 msec

MR Pause: 5.0070 msec

MCA: No

Step Size: 0.00 amu

-12-

<u>Analyte</u>	<u>Q1 Mass</u> (amu)	<u>Q3 Mass</u> (amu)	<u>Dwell</u> (msec)	<u>Parameter</u>	Start/Stop
	(umu)	(ama)	(msec)		
]					]
DP = CE = CXP =	Declustering P Collision Energy Collision Cell	gy			
Parameter Ta	able				
CUR (Curtain	Gas):		10.00		
GS1 (Gas 1):			70.00		
GS2 (Gas 2):			20.00		
IS (IonSpray V	/oltage):		-4000.00		
TEM (Temper	ature):		450.00		
ihe (Interface l	Heater):		ON		
CAD (Collisio Gas):	nally Activated	Dissociation	4.00 (nitrogen)	.·	
EP (Entrance I	Potential):		-12.00		×
Detector Para	meters (Positiv	ve)			
CEM (Channe	l Electron Multi	plier):	2600.0		
DF (Deflector)	):		200.0		

### 2.5. Preparation Of Mobile Phase

The mobile phase/diluent [2 mM ammonium acetate in 70:30 (v/v) deionized water:acetonitrile] was prepared by dissolving approximately 0.154 g of ammonium acetate in 700 mL of deionized (DI) water. ACN (300 mL) was added and the solution was stirred to achieve complete dissolution. The mobile phase was filtered through a 0.2- or 0.45-µm pore-size nylon membrane filter and vacuum-degassed. The preparation was scaled as needed.

# 2.6. PREPARATION OF THE [ ] CALIBRATION STOCK SOLUTION - 0.1 MG/ML

A calibration stock solution was prepared at a concentration of 0.1 mg [ ]/mL as follows. Approximately 0.010 g of [ ] was accurately weighed in a tared glass weigh funnel and transferred to a 100-mL volumetric flask with rinses of DI water. Additional DI water was added to yield the desired concentration, and the solution was stirred to achieve complete dissolution. The solution was stored at approximately 4°C.

# 2.7. PREPARATION OF THE | | CALIBRATION STOCK | SOLUTION - 1.0 MG/ML

A calibration stock solution was prepared at a concentration of 1.0 mg [ ]A/mL as follows. Approximately 0.100 g of [ ] was accurately weighed in a tared glass weigh funnel and transferred to a 100-mL volumetric flask with rinses of DI water. Additional DI water was added to yield the desired concentration, and the solution was stirred to achieve complete dissolution. The solution was stored at approximately 4°C.

### 2.8. Preparation Of The [ ] Quality Control Stock Solution

A quality control (QC) stock solution was prepared at a concentration of 1.0 mg [ ]/mL as follows. Approximately 0.100 g of [ ] was accurately weighed in a tared glass weigh funnel and transferred to a 100-mL volumetric flask with

rinses of DI water. Additional DI water was added to yield the desired concentration, and the solution was stirred to achieve complete dissolution. The preparation was scaled as needed. The solution was stored at approximately 4°C.

### 2.9. PREPARATION OF CALIBRATION SAMPLES

An aliquot (100  $\mu$ L) of the 1.0 mg/mL calibration stock solution was combined with 0.9 mL of DI water to prepare a 100,000 ng/mL working solution (the 0.1 mg/mL calibration stock solution was used without further dilution). Serial dilutions of the 100,000 ng/mL working or stock solution were prepared in polypropylene tubes with blank serum or urine to prepare calibration samples containing 10.0 to 1000 ng [ ]/mL. The calibration samples were processed as described in Section 2.11. (Sample Processing).

#### 2.10. Preparation Of Quality Control Samples

An aliquot (20 µL) of the QC stock solution was combined with 1.98 mL of blank rat serum, urine or cage rinse to prepare a 10,000 ng/mL sample. Serial dilutions of the 10,000 ng/mL sample were prepared in polypropylene tubes with blank serum, urine or cage rinse to prepare QC samples containing 30.0 to 750 ng [ ]/mL. Dilutional QC samples were prepared by diluting aliquots of the 10,000 ng/mL QC samples up to 20-fold with blank serum, urine or cage rinse. Dilutional QC samples were also prepared by combining an aliquot of the QC stock solution with matrix to yield QC samples at either 30,000 or 150,000 ng [ ]/mL. The 30,000 and 150,000 ng/mL QC samples were diluted up to 1000- or 5000-fold, respectively, with blank matrix. The QC samples were processed as described in Section 2.11. (Sample Processing).

### 2.11. Sample Processing

Aliquots (50  $\mu$ L) of the calibration, QC and experimental samples were transferred to 1.5-mL conical tubes. Dilutional QC samples were diluted up to 5000-fold with blank serum or urine and 50- $\mu$ L aliquots were transferred to 1.5-mL conical tubes. Experimental samples were diluted up to 5000-fold with blank serum or urine when the

initial assayed or expected concentrations were greater than 1000 ng [ ]/mL, and 50- $\mu$ L aliquots were transferred to 1.5-mL conical tubes. ACN (0.15 mL) was added to the 1.5-mL conical tubes containing the calibration, QC and experimental samples. The tubes were capped, and the samples were thoroughly mixed for a minimum of 2 minutes. The samples were centrifuged at 10,000 rpm for approximately 5 minutes. An aliquot (150  $\mu$ L) of each supernatant fraction was transferred to a 96-well plate. Samples were evaporated to dryness under a stream of nitrogen and reconstituted in 150  $\mu$ L of mobile phase.

#### 2.12. CONCENTRATION QUANTITATION

A calibration curve was constructed using Analyst[®]. The [ ] peak areas (y) and the theoretical concentrations of the calibration standards (x) were fit to a quadratic function with a  $1/x^2$  weighting, excluding the origin. Concentration and percent relative error (%RE) were calculated using Analyst[®].

#### 3. RESULTS AND DISCUSSION

#### 3.1. METHOD VALIDATION

The method for the determination of [ ] concentration in rat serum, urine and cage rinse was validated, and method sensitivity, specificity/selectivity, calibration reproducibility, ruggedness, accuracy and precision were assessed. The results of the validation are summarized in Tables 1 and 2 (Calibration Samples) and Tables 3 through 5 (QC Samples).

Under the described chromatographic conditions, the retention time for [ ] was approximately 3.0 to 3.5 minutes (Agilent 1200) or approximately 4.6 to 6.2 minutes (Symbiosis). The total run time for each analysis was approximately 7.0 minutes (Agilent 1200) or approximately 8.0 minutes (Symbiosis). Figures 1 through 35 illustrate typical chromatograms of blank samples (Figures 1 and 2), processed rat serum calibration samples (Figures 3 through 9), processed rat serum QC samples (Figures 10 through 13) and processed serum experimental samples (Figures 14 through 35).

Figures 36 through 57 illustrate typical chromatograms of blank samples (Figure 36), processed rat urine calibration samples (Figures 37 through 43), processed rat urine and cage rinse QC samples (Figures 44 through 51) and processed urine and cage rinse experimental samples (Figures 52 through 57).

#### 3.2. SENSITIVITY

The lower limit of quantitation (LLOQ) can be defined as the lowest calibration concentration that meets the validation acceptance criteria [i.e., relative standard deviation (RSD)  $\leq$  20% and percent relative error (%RE) within  $\pm$  20%]. As shown in Tables 1 and 2, the LLOQ was 10.0 ng/mL for [ ] in rat serum and urine. The inter-session (serum) and intra-session (urine) RSD and %RE values at the LLOQ are detailed in the following table.

	%RSD	%RE
Rat serum	4.4	0.82
Rat urine	6.5	1.2

#### 3.3. Specificity/Selectivity

Assay specificity/selectivity refers to the ability of the assay to specifically detect and quantitate the analyte(s) of interest from potentially interfering compounds. Assay specificity/selectivity was confirmed when assessment of the assay accuracy and precision met the acceptance criteria.

### 3.4. Ruggedness

Assay ruggedness was successfully demonstrated for this procedure because more than 1 analyst successfully performed at least 1 of the validation sessions.

### 3.5. CALIBRATION ACCEPTABILITY

During each validation session, triplicate calibration samples at 5 concentrations were prepared and analyzed. The resulting analyte peak area versus theoretical concentration data were fit to the quadratic function with  $1/x^2$  weighting, excluding the origin, using

least-squares regression analysis. The results of the regression analyses were used to back-calculate the corresponding concentrations from the peak area data. The reproducibility of the calibration curve data was considered valid when 1) the inter- or intra-session variability, expressed as %RSD, of the back-calculated concentrations at each calibration level was  $\leq 15\%$ , except at the lowest calibration level where  $\leq 20\%$  was acceptable; and 2) the mean back-calculated concentrations at each calibration level were within 15% of the theoretical values (%RE within  $\pm$  15%), except at the lowest calibration level where %RE within  $\pm$  20% was acceptable.

The back-calculated concentrations and the associated intra- and inter-session statistics for [ ] calibration samples used during the validation in rat serum and urine are shown in Tables 1 and 2. The inter-session variability of the back-calculated concentrations at each rat serum calibration level ranged from 1.2% to 4.4% RSD. The inter-session mean concentrations had %RE values ranging from -2.7% to 2.1%. The intra-session variability of the back-calculated concentrations at each rat urine calibration level ranged from 5.8% to 13% RSD. The intra-session mean concentrations had %RE values ranging from -3.3% to 4.4%. Based on the stated criteria, the reproducibility of the calibration data was acceptable.

#### 3.6. ACCURACY AND PRECISION

During each validation session, triplicate QC samples at a minimum of 3 concentrations were prepared and analyzed as described previously. Single injections were made of each processed QC sample. The results of the regression analyses were used to calculate the corresponding concentrations from the QC peak area data. The variability (RSD) of calculated QC concentration data was used as a measure of assay precision. The precision of the method was considered acceptable when the inter- or intra-session RSD of the calculated concentrations at each QC level was  $\leq$  15%. The difference between the theoretical and mean calculated QC concentrations (%RE) was used as a measure of assay accuracy. The accuracy of the method was considered acceptable when

the inter- or intra-session mean calculated concentration at each QC level had a %RE value within  $\pm$  15%.

The calculated concentrations and the associated intra- and inter-session statistics for the [ ] assay QC samples used during the validation in rat serum, urine and cage rinse are summarized in Tables 3 through 5. The inter-session variability and the %RE values for the inter- (serum) or intra- (urine and cage rinse) session mean concentrations are summarized in the following table.

	%RSD	<u>%RE</u>
Rat serum	4.4 to 6.3	-4.9 to 3.6
Rat urine	6.5 to 14	-6.7 to 11
Rat cage rinse	1.8 to 4.3	-6.7 to 3.3

Based on the stated criteria, the precision and accuracy of the [ ] assays were acceptable.

#### 3.7. STABILITY

Stability of [ ] was evaluated in stock solutions and in samples after short-term (4 hours) room temperature storage, during the freeze-thaw process and during frozen storage. According to the protocol, stability was verified if the mean measured post-storage (or treatment) analyte concentration was not less than 85% of the corresponding time-zero concentration.

### 3.7.1. STABILITY OF [ ] IN REFRIGERATED STOCK SOLUTIONS

Stability of [ ] in stock solutions was evaluated. Standards at 400 ng/mL in water were prepared from stock solutions after 11 days of refrigerated storage. The [ ] peak areas of the standards prepared after storage of the stock solution were compared to standards prepared from the freshly prepared (day 0) standard [ ] peak areas.

The mean areas of the standards prepared from stock solutions after 11 days of refrigerated storage were 99.1% of the mean time-zero area (Table 6). Therefore, [ ] in stock solutions was considered to be stable for at least 11 days of refrigerated storage.

# 3.7.2. Long Term Frozen Stability Of [ ] In Serum And Urine Samples

Rat serum and urine samples were fortified at 30.0 and 750 ng [ ]/mL, analyzed in at least duplicate for time-zero concentration, stored frozen at approximately -20°C for up to 127 days and reanalyzed in triplicate to evaluate test article stability.

The mean measured concentrations in the rat serum samples after frozen storage of up to 124 days ranged from 94.3% to 115% of the corresponding time-zero concentrations (Table 7), which met the specified criteria. The mean measured concentrations in the rat urine samples after frozen storage of up to 127 days ranged from 107% to 113% of the corresponding time-zero concentrations (Table 8), which met the specified criteria. Therefore, [ ] in rat serum and urine samples was considered to be stable for up to 124 and 127 days of frozen (-70°C) storage, respectively.

# 3.7.3. STABILITY OF [ ] IN SERUM AND URINE SAMPLES AT ROOM TEMPERATURE

Blank rat serum and urine were fortified at 30.0 and 750 ng [ ]/mL, analyzed in triplicate to establish time-zero concentration, stored frozen at approximately -20°C and allowed to thaw at room temperature. After approximately 4 hours of room temperature storage, the samples were reanalyzed in triplicate to assess test article stability.

The mean measured concentrations in the 30.0 and 750 ng/mL serum samples after approximately 4 hours of room temperature storage were 110% and 94.7% of the corresponding time-zero concentrations (Table 9), which met the specified criteria for stability. The mean measured concentrations in the 30.0 and 750 ng/mL urine samples after approximately 4 hours of room temperature storage were 108% and 106% of the corresponding time-zero concentrations (Table 10), which met the specified criteria for

stability. Therefore, [ ] in rat serum and urine samples was considered to be stable for at least 4 hours of room temperature storage.

#### 3.7.4. Freeze-Thaw Stability Of [ ] IN SERUM AND URINE SAMPLES

Rat serum and urine samples were fortified at 30.0 and 750 ng [ ]/mL. Triplicate samples from each concentration level were used to evaluate the stability of the analyte after each of a minimum of 3 freeze-thaw cycles. The samples were frozen at approximately -20°C and thawed (1 cycle), and the process repeated a minimum of 2 more times (cycles 2 and 3) for the analysis of freeze-thaw stability.

The mean measured concentrations in rat serum samples after up to 3 freeze-thaw cycles ranged from 96.3% to 120% of the corresponding time-zero concentrations (Table 11), which met the specified acceptance criteria. The mean measured concentrations in rat urine samples after up to 6 freeze-thaw cycles ranged from 109% to 113% of the corresponding time-zero concentrations (Table 12), which met the specified acceptance criteria. Therefore, [ ] in rat serum and urine samples was considered to be stable through at least 3 and 6 freeze-thaw cycles, respectively.

#### 3.8. ANALYSIS OF EXPERIMENTAL SAMPLES

Rat serum and urine samples from study day 0 were analyzed for [ ] concentration. The results are summarized in Tables 13 and 14.

In addition to the experimental samples, each set of analyses consisted of at least duplicate calibration samples at 5 concentrations, 1 solvent blank, 3 blank matrix samples and at least triplicate QC samples at a minimum of 3 concentrations. Dilutional QC samples were analyzed as needed. For an analytical run to be considered valid, at least two-thirds of the QC samples (with at least 1 at each concentration level) could not deviate more than  $\pm$  15% from the QC target concentrations. Tables 15 and 16 summarize the results of the calibration and QC samples. All reported results are from analyses that met the acceptance criteria.

#### 4. CONCLUSION

In this study, the method for the analysis of [ ] concentration in rat serum, urine and cage rinse was validated over the concentration range of 10.0 to 1000 ng [ ]/mL of matrix using a 50- $\mu$ L sample. Analyte stability in stock solutions and in stored rat serum and urine samples was assessed. [ ] in stock solutions was stable for at least 11 days of refrigerated storage. [ ] in rat serum was stable through at least 3 freeze-thaw cycles, for at least 4 hours of room temperature storage and for at least 124 days of frozen (approximately -20°C) storage. [ ] in rat urine was stable through at least 6 freeze-thaw cycles, for at least 4 hours of room temperature storage and for at least 127 days of frozen (approximately -20°C) storage. Analysis of experimental serum samples resulted in levels ranging from <LLOQ to 100,720 ng [ ]/mL. Analysis of experimental urine samples resulted in total [ ] values ranging from 4.44 to 898  $\mu$ g. Analysis of experimental cage rinse samples resulted in total [ ] values ranging from 1.87 to 1006  $\mu$ g.

**TABLES 1 - 16** 

Table 1: [ ] Back-Calculated Concentrations And Intra- And Inter-Session Statistics Of Calibration Samples In Rat Serum

Theo. Conc. (ng/mL)	10.0	30.0	100	300	1000
Session 1 (28 Feb 200	7), <i>18-53</i>	4006a, and	lyst DKF	•	
Sample 1	10.4	28.5	104	301	1015
Sample 2	10.4	28.8	105	296	990
Sample 3	9.41	29.0	105	286	1003
Intra-session Statistics					
n	3	3	3	3	3
Mean	10.1	28.8	105	294	1003
· SD	0.60	0.26	0.77	7.7	13
RSD	5.9	0.90	0.73	2.6	1.3
%RE	0.93	-4.0	4.8	-1.9	0.26
Session 2 (28 Feb 200	7), 18-53	4006b, and	lyst DKF	)	
Sample 1	9.95	**	97.9	318	974
Sample 2	10.3	29.9	98.3	305	982
Sample 3	9.76	30.0	98.6	299	1020
Intra-session Statistics					
n	3	2	3	3	3
Mean	10.0	29.9	98.3	307	992
SD	0.29	0.10	0.36	9.6	24
RSD	2.9	0.32	0.37	3.1	2.4
%RE	0.14	-0.17	-1.7	2.4	-0.79

^{**} Data point not included in calculation of statistics based on test for outliers.

Table 1: [ ] Back-Calculated Concentrations And Intra- And Inter-Session Statistics Of Calibration Samples In Rat Serum

Theo. Conc. (ng/mL)	10.0	30.0	100	300	1000		
Session 3 (13 Mar 200	Session 3 (13 Mar 2007), 18-534006g, analyst SMH						
Sample 1	10.1	27.3	103	322	1007		
Sample 2	10.1	29.6	99.1	308	1000		
Sample 3	10.4	26.7	104	293	981		
Intra-session Statistics							
n	. 3	3	3	3	3		
Mean	10.2	27.9	102	308	996		
SD	0.16	1.5	2.5	14	14		
RSD	1.6	5.5	2.5	4.6	1.4		
%RE	2.1	-7.1	1.9	2.6	-0.42		
Session 4 (13 Mar 200	77), I8-53	4006h, an	alyst SM	H			
Sample 1	10.2	29.6	98.9	302	1001		
Sample 2	10.2	30.5	98.2	303	995		
Sample 3	9.68	28.9	99.8	309	997		
Intra-session Statistics							
n	3	3	3	3	3		
Mean	10.0	29.7	99.0	305	998		
SD	0.31	0.81	0.78	3.8	3.2		
RSD	3.1	2.7	0.79	1.2	0.32		
%RE	0.42	-1.1	-1.0	1.6	-0.22		

Table 1: [ ] Back-Calculated Concentrations And Intra- And Inter-Session Statistics Of Calibration Samples In Rat Serum

Theo. Conc. (ng/mL)	10.0	30.0	100	300	1000
Session 5 (13 Mar 200	7), I8-53	4006i, ana	lyst HLC		
Sample 1	10.0	28.8	94.0	320	1003
Sample 2	10.8	29.1	93.8	322	979
Sample 3	9.29	32.4	98.1	295	1003
Intra-session Statistics	<u></u> '				
n	3	3	3	3	3
Mean	10.0	30.1	95.3	312	995
SD	0.73	2.0	2.4	15	14
RSD	7.3	6.6	2.5	4.8	1.4
%RE	0.24	0.28	<del>-</del> 4.7	4.1	-0.51
Session 6 (14 Mar 200	07), I8-53	4006j, and	lyst HLC		
Sample 1	9.96	29.7	98.7	311	1002
Sample 2	9.44	28.4	98.8	320	998
Sample 3	10.9	29.2	97.3	302	984
Intra-session Statistics	'				
n	3	3	3	3	3
Mean	10.1	29.1	98.3	311	995
SD	0.75	0.67	0.83	9.2	10
RSD	7.5	2.3	0.85	2.9	0.96
%RE	1.0	-3.0	-1.7	3.6	-0.52
Inter-session Statistics					
n	18	17	18	18	18
Mean	10.1	29.2	99.6	306	996
SD	0.45	1.2	3.4	11	12
RSD	4.4	4.3	3.4	3.5	1.2
%RE	0.82	-2.7	-0.42	2.1	-0.37

Table 2: [ ] Back-Calculated Concentrations And Intra-Session Statistics Of Calibration Samples In Rat Urine

				·		
10.0	30.0	100	300	1000		
Session 1 (15 Mar 2007), 18-534006l, analyst HLC						
9.36	29.2	110	328	1035		
10.6	32.8	84.0	283	1017		
10.4	25.1	99.0	328	927		
	···					
3	3	3	3	3		
10.1	29.0	97.5	313	993		
0.66	3.9	13	26	58		
6.5	13	13	8.3	5.8		
1.2	-3.3	-2.5	4.4	-0.71		
	9.36 10.6 10.4 3 10.1 0.66 6.5	9.36 29.2 10.6 32.8 10.4 25.1 3 3 10.1 29.0 0.66 3.9 6.5 13	9.36 29.2 110 10.6 32.8 84.0 10.4 25.1 99.0 3 3 3 3 10.1 29.0 97.5 0.66 3.9 13 6.5 13 13	77), 18-534006l, analyst HLC       9.36     29.2     110     328       10.6     32.8     84.0     283       10.4     25.1     99.0     328       3     3     3     3       10.1     29.0     97.5     313       0.66     3.9     13     26       6.5     13     13     8.3		

534006.xls Cal (urine) Printed: 09/19/07 12:58 PM

Table 3: [ ] Concentrations And Intra- And Inter-Session Statistics Of Quality Control Samples In Rat Serum

Theo. Conc. (ng/mL)	30.0	250	750	10000
Session 1 (28 Feb 200	07), I8-53	4006a, an	alyst DKI	P
Sample 1	28.1	238	714	9425
Sample 2	28.7	241	723	9384
Sample 3	29.2	244	723	9490
Intra-session Statistics	7 .			
n	3	3	3	3
Mean	28.6	241	720	9433
SD	0.54	3.2	5.4	54
RSD	1.9	1.3	0.75	0.57
%RE	-4.5	-3.6	-4.0	-5.7
Session 2 (28 Feb 200	77), 18-53	4006b, an	alyst DKI	D
Sample 1	31.3	239	748	9709
Sample 2	29.2	241	754	9685
Sample 3	28.1	248	747	9626
Intra-session Statistics	3			
n	3	3	3	3
Mean	29.5	243	750	9673
SD	1.6	4.6	3.8	43
RSD	5.4	1.9	0.51	0.44
%RE	-1.6	-3.0	-0.059	-3.3

534006.xls QC (serum) Printed: 09/19/07 12:58 PM

Table 3: [ ] Concentrations And Intra- And Inter-Session Statistics Of Quality Control Samples In Rat Serum

Theo. Conc. (ng/mL)	30.0	250	750	10000
Session 3 (13 Mar 200	07), I8-53	4006g, an	alyst SM	H
Sample 1	30.3	268	798	***
Sample 2	26.2	268	773	***
Sample 3	32.1	274	786	***
Intra-session Statistics	Ţ		• • • • • • • • • • • • • • • • • • • •	
n	3	3	3	na
Mean	29.6	270	786	na
SD	3.0	3.3	12	na
RSD	10.3	1.2	1.6	na
%RE	-1.5	8.0	4.8	na
Session 4 (13 Mar 200	07), I8-53	4006h, ar	alyst SM	H
Sample 1	26.7	256	805	11265
Sample 2	27.5	260	815	10601
Sample 3	30.3	265	799	10886
Intra-session Statistics	7		•	
n	3	3	3	3
Mean	28.2	260	806	10917
SD	1.9	4.4	7.9	333
RSD	6.8	1.7	0.99	3.1
%RE	-6.2	4.1	7.5	9.2

na = not applicable

534006.xls QC (serum) Printed: 09/19/07 12:58 PM

^{***} QC sample data not included due to preparation error.

Table 3: [ ] Concentrations And Intra- And Inter-Session Statistics Of Quality Control Samples In Rat Serum

Theo. Conc. (ng/mL)	30.0	250	750	10000		
Session 5 (13 Mar 200	7), I8-53	4006i, and	alyst HLC	<b>Y</b>		
Sample 1	27.7	245	788	10142		
Sample 2	25.2	247	807	10242		
Sample 3	28.9	271	826	10656		
Intra-session Statistics	3					
n	3	3	3	3		
Mean	27.2	254	807	10347		
SD	1.9	15	19	273		
RSD	7.0	5.8	2.4	2.6		
%RE	-9.2	1.8	7.5	3.5		
Session 6 (14 Mar 20)	07), 5340	06j, analy	st HLC			
Sample 1	28.7	246	807	***		
Sample 2	28.5	246	772	***		
Sample 3	26.8	249	802	***		
Intra-session Statistics	3					
n	3	3	3	na		
Mean	28.0	247	794	na		
SD	1.1	1.7	19	na		
RSD	3.8	0.69	2.4	na		
%RE	-6.7	-1.2	5.8	na		
Inter-session Statistics	Inter-session Statistics					
n	18	18	18	12		
Mean	28.5	253	777	10093		
SD	1.8	12	34	636		
RSD	6.2	4.8	4.4	6.3		
%RE	-4.9	1.0	3.6	0.93		

^{***} QC sample data not included due to preparation error.

na = not applicable

534006.xls QC (serum)

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Table 4: [ ] Concentrations And Intra-Session Statistics Of Quality Control Samples In Rat Urine

Theo. Conc. (ng/mL)	30.0	250	750	10000		
Session 1 (15 Mar 2007), 18-534006l, analyst HLC						
Sample 1	27.9	242	793	9978		
Sample 2	36.6	216	903	12540		
Sample 3	34.7	242	801	10064		
Intra-session Statistics	,					
n	3	3	3	3		
Mean	33.1	233	832	10861		
SD	4.6	15	62	1455		
RSD	14	6.5	7.4	13		
%RE	10	-6.7	11	8.6		

534006.xls QC (urine) Printed: 09/19/07 12:58 PM

Table 5: [ ] Concentrations And Intra-Session Statistics Of Quality Control Samples In Cage Rinse

		<b>-</b>	_					
Theo. Conc. (ng/mL)	30.0	250	750	30000				
Session 1 (28 Mar 2007), 16-53400611, analyst SMH								
Sample 1	31.2	240	750	27523				
Sample 2	29.4	232	783	28519				
Sample 3	29.9	253	783	27956				
Intra-session Statistics	,							
n	3	3	3	3				
Mean	30.2	242	772	27999				
SD	0.91	11	19	499				
RSD	3.0	4.3	2.4	1.8				
%RE	0.56	-3.3	3.0	-6.7				

534006.xls QC (cage rinse) Printed: 09/19/07 12:58 PM

# PHARMACOKINETIC (IN BLOOD) <u>AND EXCRETION STUDY OF [ ] IN RATS</u> Table 6: Refrigerated (4°C) Stability Of [ ] In DI Water Stock Solutions

Storage <u>Duration</u> (days)	<u>Run #</u> (534006-)	<u>Ref. #</u> (534006-)	<u>Area</u>	Mean <u>Area</u>	<u>RSD</u> (%)	% of Time Zero
O ^a	I6-1109 I6-1110	201-6 201-8	26400000 26600000	26500000	0.53	NA
11 ^a	I6-1107 I6-1108	201-2 201-4	26400000 26100000	26250000	0.81	99.1

a. Results for Time Zero and 11-day were in Sequence I6-534006j

NA = not applicable

534006.xls Stock Printed: 19Sep2007 12:58 PM

Table 7: Stability of [ ] In Rat Serum - Long-Term Frozen (-20°C) Storage

Theo. Conc. (ng/mL)	Storage <u>Duration</u> (Days)	<u>Run#</u> (534006-)	<u>Ref. #</u> (534006-)	Assay Conc. (ng/mL)	Mean <u>Conc.</u> (ng/mL)	<u>SD</u>	<u>RSD</u> (%)	% of Theo.	% of <u>Time Zero</u>
30.0	$0^a$	I6-0684	155-2	28.0	27.7	0.46	1.7	92.3	na
	0 ^a	I6-0723	155-3	27.4					
	11 ^b	I6-1155	209-1	30.5	30.8	0.58	1.9	103	111
	11 ^b	I6-1156	209-2	31.5					
	11 ^b	I6-1157	209-3	30.5					
	124 ^c	I6-1675	264-1	31.0	31.9	0.86	2.7	106	115
	124°	I6-1676	264-2	32.2					
	124 ^c	I6-1677	264-3	32.6					
750		I6-0686	155 0	922	806	24	2.0	107	
/50	0	I6-0686 I6-0725	155-8 155-9	823 789	800	24	2.9	107	na
	11	I6-1158	209-4	773	765	6.7	0.87	102	95.0
	11	I6-1159	209-5	760					
	11	I6-1160	209-6	764					
	124	I6-1678	264-4	772	760	11	1.5	101	94.3
	124	I6-1679	264-5	750					
	124	I6-1680	264-6	759					

a. Results for 0 day were in Sequence I6-534006f1

na= not applicable

534006.xls LT (serum) Printed: 09/24/07 7:38 AM

b. Results for 11 days were in Sequence I6-534006k

c. Results for 124 days were in Sequence I6-534006p1

Table 8: Stability of [ ] In Rat Urine - Long-Term Frozen (-20°C) Storage

Theo. Conc. (ng/mL)	Storage Duration (Days)		<u>Run#</u> (534006-)	<u>Ref. #</u> (534006-)	Assay Conc. (ng/mL)	Mean <u>Conc.</u> (ng/mL)	<u>SD</u>	<u>RSD</u> (%)	% of Theo.	% of Time Zero
30.0	0	a	I6- 0751	165-1	27.4	27.1	1.3	4.8	90.5	na
	0	a	I6-0 780	165-2	25.8					
	0	a	16-0 804	165-3	28.3					
	14	b	I6-1411	235-1	31.6	30.3	1.2	3.8	101	112
	14	Ъ	I6-1412	235-2	30.0					
	14	Ъ	I6-1413	235-3	29.3					
	127	c	I6-1800	273-1	31.4	28.9	2.9	10	96.5	107
	127	С	I6-1801	273-2	25.7					
	127	С	I6-1802	273-3	29.8	-				
750	0	а	I6-0753	165-7	697	743	41	5.5	99.0	na
	0	a	I6-0782	165-8	775					
	0 .	a	16-0806	165-9	755					
	14	ъ	I6-1420	235-10	809	839	36	4.2	112	113
	14	b	I6-1421	235-11	878					
	14	Ъ	I6-1422	235-12	831					
	127	c	I6-1803	273-4	782	836	50	6.0	111	113
	127	c	I6-1804	273-5	845					
	127	С	I6-1805	273-6	881					

a. Results for 0 day were in Sequence I6-534006g

na= not applicable

534006.xls LT (urine) Printed: 09/19/07 12:58 PM

b. Results for 14 days were in Sequence I6-534006m

c. Results for 127 days were in Sequence I6-534006q1

# PHARMACOKINETIC (IN BLOOD) AND EXCRETION STUDY OF [ ] IN RATS Table 9: Room Temperature Stability Of [ ] In Rat Serum

Theo. Conc. (ng/mL)	Storage <u>Duration</u> (hours)	<u>Run #</u> (534006-)	<u>Ref. #</u> (534006-)	Analyzed Conc (ng/mL)	% Target	Mean <u>Conc</u> (ng/mL)	RSD (%)	Mean <u>% Target</u>	% of Time <u>Zero</u>
30.0	$0^a$	I6-0684 I6-0723	155-2 155-3	28.0 27.4	93.4 91.2	27.7	1.7	92.3	NA
(apṛ	4 ^b proximately)	I6-1175 I6-1176 I6-1177	210-1 210-2 210-3	30.2 30.4 30.6	101 101 102	30.4	0.56	101	110
750	0 ^a	I6-0686 I6-0725	155-8 155-9	823 789	110 105	806	2.9	107	NA
(app	4 ^b proximately)	I6-1178 I6-1179 I6-1180	210-4 210-5 210-6	763 772 755	102 103 101	763	1.1	102	94.7

NA = not applicable

534006.xls ST (serum) Printed: 19Sep2007 12:58 PM

a. Results for Time Zero were in Sequence I6-534006f1

b. Results for 4-hour were in Sequence I6-534006k

# PHARMACOKINETIC (IN BLOOD) <u>AND EXCRETION STUDY OF [ ] IN RATS</u> Table 10: Room Temperature Stability Of [ ] In Rat Urine

Theo. Conc. (ng/mL)	Storage <u>Duration</u> (hours)	Run # (534006-)	<u>Ref. #</u> (534006-)	Analyzed <u>Conc</u> (ng/mL)	% <u>Target</u>	Mean <u>Conc</u> (ng/mL)	RSD (%)	Mean <u>% Target</u>	% of Time Zero
30.0	$0^{a}$	I6- 0751 I6-0780 I6-0804	165-1 165-2 165-3	27.4 25.8 28.3	91.3 85.8 94.3	27.1	4.8	90.5	NA
(app	4 ^b proximately)	I6-1391 I6-1392 I6-1393	233-1 233-2 233-3	31.2 28.6 28.3	104 95.4 94.3	29.4	5.5	98.0	108
750	$0^a$	I6-0753 I6-0780 I6-0806	165-7 165-8 165-9	697 775 755	92.9 103 101	743	5.5	99.0	NA
(app	4 ^b proximately)	I6-1394 I6-1395 I6-1396	233-4 233-5 233-6	761 766 826	102 102 110	785	4.6	105	106

NA = not applicable

534006.xls ST (urine) Printed: 19Sep2007 12:58 PM

a. Results for Time Zero were in Sequence I6-534006g

b. Results for 4-hour were in Sequence I6-534006m

Table 11: Freeze-Thaw Stability Of [ ] In Rat Serum

Theo. Conc. (ng/mL)	# of Cycles	Run # (534006-)	<u>Ref. #</u> (534006-)	Analyzed Conc (ng/mL)	% Target	Mean <u>Conc</u> (ng/mL)	RSD (%)	Mean <u>% Target</u>	% of Time Zero
30.0	0 ^a	I6-0684 I6-0723	155-2 155-3	28.0 27.4	93.4 91.2	27.7	1.7	92.3	NA
•	1 ^b	I6-1694 I6-1695 I6-1696	265-1 265-2 265-3	31.9 32.1 32.6	106 107 109	32.2	1.2	107	116
	2	I6-1697 I6-1698 I6-1699	265-4 265-5 265-6	32.7 33.4 32.9	109 111 110	33.0	1.0	110	119
	3	I6-1700 I6-1701 I6-1702	265-7 265-8 265-9	33.9 33.0 32.6	113 110 109	33.2	2.1	111	120
750	0ª	I6-0686 I6-0725	155-8 155-9	823 789	110 105	806	2.9	107	NA
	1 ^b	I6-1703 I6-1704 I6-1705	265-10 265-11 265-12	784 780 770	105 104 103	778	0.92	104	96.6
	2	I6-1706 I6-1707 I6-1708	265-13 265-14 265-15	768 765 795	102 102 106	776	2.2	103	96.3
	3	I6-1709 I6-1710 I6-1711	265-16 265-17 265-18	790 792 794	105 106 106	792	0.26	106	98.2

a. Results for Time Zero were in Sequence I6-534006f1

534006.xls F-T (serum)

NA = not applicable Printed: 19Sep2007 12:58 PM

b. Results for cycles 1 through 3 were in Sequence I6-534006p1

# PHARMACOKINETIC (IN BLOOD) <u>AND EXCRETION STUDY OF [ ] IN RATS</u> Table 12: Freeze-Thaw Stability Of [ ] In Rat Urine

Theo. Conc. (ng/mL)	# of Cycles	<u>Run #</u> (534006-)	<u>Ref. #</u> (534006-)		% Target	Mean <u>Conc</u> (ng/mL)	RSD (%)		% of Time Zero
30.0	0 ^a	I6- 0751 I6-0780 I6-0804	165-1 165-2 165-3	27.4 25.8 28.3	91.3 85.8 94.3	27.1	4.8	90.5	NA
	1 ^b	I6-1411 I6-1412 I6-1413	235-1 235-2 235-3	31.6 30.0 29.3	105 100 97.6	30.3	3.8	101	112
	2	I6-1414 I6-1415 I6-1416	235-4 235-5 235-6	31.2 28.1 30.0	104 93.6 99.9	29.8	5.3	99.2	110
	3	I6-1417 I6-1418 I6-1419	235-7 235-8 235-9	30.7 28.5 29.4	102 95.0 98.0	29.5	3.7	98.4	109
	6°	I6-1819 I6-1820 I6-1821	274-1 274-2 274-3	32.9 29.9 29.5	110 99.6 98.2	30.8	6.2	103	113

NA = not applicable

a. Results for cycle 0 were in Sequence I6-534006g

b. Results for cycles 1 to 3 were in Sequence I6-534006m

c. Results for cycle 6 were in Sequence I6-534006q1

Table 12: Freeze-Thaw Stability Of [ ] In Rat Urine (Continued)

Theo. Conc. (ng/mL)	# of Cvcles	<u>Run #</u> (534006-)	<u>Ref. #</u> (534006-)	Analyzed Conc (ng/mL)	% Target	Mean <u>Conc</u> (ng/mL)	RSD (%)	Mean <u>% Target</u>	% of Time <u>Zero</u>
750	0 ^a	I6-0753	165-7	697	92.9	743	5.5	99.0	NA
		I6-0782	165-8	775	103				
		I6-0806	165-9	755	101				
	1 ^b	I6-1420	235-10	809	108	839	4.2	112	113
	•	I6-1421	235-11	878	117	037	1,2	112	115
		I6-1422	235-12	831	111				
	2	I6-1423	235-13	861	115	810	15	108	109
	_	I6-1424	235-14	899	120	010	10	100	107
		I6-1425	235-15	668	89.1				
	3	I6-1426	235-16	823	110	817	6.8	109	110
	-	I6-1427	235-17	759	101	91,	•••	10)	110
		I6-1428	235-18	870	116				
	6°	I6-1822	274-4	830	111	825	6.5	110	111
		I6-1823	274-5	770	103				
		I6-1824	274-6	876	117				

NA = not applicable

534006.xls F-T (urine) Printed: 19Sep2007 12:58 PM

a. Results for cycle 0 were in Sequence I6-534006g

b. Results for cycles 1 to 3 were in Sequence I6-534006m

c. Results for cycle 6 were in Sequence I6-534006q1

Table 13: Rat Serum Experimental Sample [ ] Concentration Data

			<b></b>			[ ]	
<u>Ref. #</u>	Animal No.	<u>Day</u>	<u>Group</u>	<u>Sex</u>	<u>Timepoint</u>	Conc.	
(534006-)					(hrs)	(ng/mL)	
195-1	46662	0	1	M	. 0	<lloq< td=""><td>1</td></lloq<>	1
195-2	46663	0	1	M	0	<lloq< td=""><td></td></lloq<>	
195-3	46665	0	1	M	0	<lloq< td=""><td></td></lloq<>	
195-4	46678	0	1	F	0	<lloq< td=""><td></td></lloq<>	
195-5	46679	0	1	F	0	<lloq< td=""><td></td></lloq<>	
195-6	46680	0	1	F	0	<lloq< td=""><td></td></lloq<>	
196-1	46666	0 .	1	M	0.0333	85509	
196-2	46669	0	1	M	0.0333	16736	
196-3	46672	0	1	M	0.0333	100720	
196-4	46681	0	1	F	0.0333	88738	
196-5	46684	0	1	F	0.0333	98125	
196-6	46685	0	1	F	0.0333	97362	
196-7	46673	0	1	M	0.167	71736	
196-8	46674	0	1	M	0.167	59438	
196-9	46676	0	1	M	0.167	76212	
196-10	46686	0	1	F	0.167	53064	
196-11	46688	0	1	F	0.167	43359	
196-12	46691	0	1	F	0.167	54242	
196-13	46662	0	1	M	0.333	65371	
196-14	46663	0	1	M	0.333	55817	
196-15	46665	0	1	M	0.333	65870	
196-16	46678	0	1	F	0.333	30804	
196-17	46679	0	1	F	0.333	27712	
196-18	46680	0	1	F	0.333	31506	
197-1	46666	0	1	M	0.5	51702	
197-2	46669	0	1	M	0.5	19975	
197-3	46672	0	1	M	0.5	57560	
197-4	46681	0	1	F	0.5	27329	
197-5	46684	0	1	F	0.5	24091	
197-6	46685	0	1	F	0.5	19724	
					•		

^{1. &}lt; LLOQ = not detected or less than the lower limit of quantitation (10 ng/mL)

534006.xls Summary (serum) Printed: 09/19/07 12:58 PM

Table 13: Rat Serum Experimental Sample [ ] Concentration Data

	-					
	_	_	~			
Animal No.	<u>Day</u>	Group	<u>Sex</u>			
16650	•	_				
		_				
46679						
46680			F		625	
46666	0	1	M	5	8627	
46669	. 0	1	M	5	20677	
46672		1	M	5	17947	
46681	0	1	F	5	639	
46684	0	1	F	- 5	285	
46685	0	1	F	5	160	
46673	0	1	M	7	24227	
46674	0	1	M	7	7111	
46676	0	1	M	. 7	20778	
46686	0	1	F	7	409	
46688	0	1 .	F	7	77.0	
46691	0	1	F	7	176	
46662	0	1	M	24	395	
46663	0	1	M	24	73.2	
46665	0 -	1	M	24	106	
46678	0	1	F	24	13.2	
46679	0	1	F	24	9.90	
46680	0	1	F	24	12.5	
46666	0	1	M	48	18.3	
46669	0	1	M	48	178	
46672	0	1	M	48	23.0	
46681	0	1	F	48	13.4	
46684	0	1	F	48	<lloq< td=""><td>1</td></lloq<>	1
46685	0	1	F	48	13.5	
	46666 46669 46672 46681 46684 46685 46673 46676 46686 46688 46691 46662 46663 46665 46678 46679 46680 46666 46669 46672 46681 46684	46673 0 46674 0 46676 0 46686 0 46688 0 46661 0 46662 0 46663 0 46665 0 46679 0 46669 0 46672 0 46681 0 46685 0 46674 0 46686 0 46686 0 46669 0 46674 0 46686 0 46687 0 46688 0 46669 0 46679 0 46670 0 46688 0 46669 0 46670 0 46680 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46661 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46661 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0 46660 0	46673       0       1         46674       0       1         46676       0       1         46686       0       1         46688       0       1         46688       0       1         46691       0       1         46662       0       1         46663       0       1         46665       0       1         46679       0       1         46680       0       1         46669       0       1         46672       0       1         46681       0       1         46682       0       1         46683       0       1         46674       0       1         46686       0       1         46687       0       1         46688       0       1         46662       0       1         46679       0       1         46679       0       1         46679       0       1         46679       0       1         46679       0       1         46680	46673       0       1       M         46674       0       1       M         46676       0       1       M         46686       0       1       F         46688       0       1       F         46689       0       1       F         46691       0       1       M         46662       0       1       M         46663       0       1       M         46665       0       1       M         46678       0       1       F         46679       0       1       F         46680       0       1       M         46672       0       1       M         46681       0       1       F         46682       0       1       F         46683       0       1       F         46684       0       1       F         46685       0       1       M         46674       0       1       M         46676       0       1       M         46686       0       1       F         46681	(hrs)           46673         0         1         M         1           46674         0         1         M         1           46676         0         1         M         1           46676         0         1         M         1           46686         0         1         F         1           46688         0         1         F         1           46689         0         1         F         1           46662         0         1         M         3           46663         0         1         M         3           46665         0         1         M         3           46678         0         1         F         3           46679         0         1         F         3           46680         0         1         M         5           46669         0         1         M         5           46681         0         1         F         5           46674         0         1         M         7           46685         0         1         F         7 </td <td>  (hrs) (ng/mL)   46673</td>	(hrs) (ng/mL)   46673

^{1. &}lt; LLOQ = not detected or less than the lower limit of quantitation (10 ng/mL)

534006.xls Summary (serum) Printed: 09/19/07 12:58 PM

# Table 14: Rat Urine/Cage Rinse Experimental Sample [ ] Concentration Data

		•			Urine/	
				[ ]	Cage Rir	ise Total
<u>Ref. #</u>	Animal No.	<b>Timepoint</b>	<u>Matrix</u>	Conc.	Volume	<u>[_].</u>
(534006-)				(ng/mL)	(mL)	(μg)
221-1	46664	0-6 hr	urine	345340	2.6	898
221-4	46670	0-6 hr	urine	394510	1.4	552
221-5	46682	0-6 hr	urine	564250	1.3	734
230-1	46683	0-6 hr	urine	1533800	0.3	460
221-8	46690	0-6 hr	urine	394380	2.0	789
231-1	46664	0-6 hr	cage rinse	10363	11	114
221-3	46667	0-6 hr	cage rinse	96774	10.4	1006
222-1	46670	0-6 hr	cage rinse	4828	13.2	63.7
222-2	46682	0-6 hr	cage rinse	8214	10.6	87.1
221-7	46683	0-6 hr	cage rinse	12178	13.4	163
222-3	46690	0-6 hr	cage rinse	7205	11.8	85.0
221-9	46664	6-12 hr	urine	141640	4.4	623
221-10	46667	6-12 hr	urine	104600	4.6	481
221-11	46670	6-12 hr	urine	89395	5.6	501
253-1	46682	6-12 hr	urine	12069	4.4	53.1
221-13	46683	6-12 hr	urine	74686	4.2	314
221-14	46690	6-12 hr	urine	113540	4.0	454
222-4	46664	6-12 hr	cage rinse	3614	14.0	50.6
222-5	46667	6-12 hr	cage rinse	1849	14.0	25.9
222-6	46670	6-12 hr	cage rinse	2295	14.0	32.1
167-19	46682	6-12 hr	cage rinse	671	14.8	9.94
222-7	46683	6-12 hr	cage rinse	2118	17.0	36.0
222-8	46690	6-12 hr	cage rinse	2038	15.8	32.2
221-15	46664	12-24 hr	urine	8082	12.6	102
221-16	46667	12-24 hr	urine	15157	9.0	136
221-17	46670	12-24 hr	urine	8782	10.8	94.8
254-1	46682	12-24 hr	urine	1200	7.2	8.64
232-2	46683	12-24 hr	urine	1315	6.2	8.16
168-11	46690	12-24 hr	urine	616	7.2	4.44
168-2	46664	12-24 hr	cage rinse	753	14.0	10.5
222-9	46667	12-24 hr	cage rinse	948	7.8	7.40
168-6	46670	12-24 hr	cage rinse	632	9.8	6.20
168-8	46682	12-24 hr	cage rinse	238	10.9	2.59
168-10	46683	12-24 hr	cage rinse	342	10.8	3.70
168-12	46690	12-24 hr	cage rinse	123	15.2	1.87

534006.xls Summary (urine) Printed: 09/19/07 12:58 PM

Table 15: [ ] Calibration Samples - Routine Analyses							
<u>Sequence</u>	<u>QC</u>	<u>Run #</u>	Ref#	Target Conc	Analyzed Conc	<u>% RE</u>	
		(534006-)	(534006-)	(ng/mL)	(ng/mL)		
I6-534006f1	Cal 1	I6-0678	154-2	10.0	10.0	0.20	
serum		I6-0728	154-3	10.0	9.82	-1.9	
	Cal 2	I6 <b>-</b> 0679	154-5	30.0	29.8	-0.82	
		16-0729	154-6	30.0	31.2	4.1	
	Cal 3	16-0680	154-8	100	107	7.2	
		I6-0730	154-9	100	104	4.3	
	Cal 4	I6 <b>-</b> 0681	154-11	300	288	-3.9	
		I6-0731	154-12	300	252	-16	
	Cal 5	16-0682	154-14	1000	1103	10	
		I6-0732	154-15	1000	1066	6.6	

	Table 1	Table 15: [ ] Calibration Samples - Routine Analyses					
Sequence	<u>QC</u>	<u>Run #</u>	Ref#	Target Conc	Analyzed Conc	<u>% RE</u>	
		(534006-)	(534006-)	(ng/mL)	(ng/mL)		
I6-534006g	Cal 1	I6-0745	164-1	10.0	11.6	16	
urine		I6-0774	164-2	10.0	10.6	6.0	
		I6-0809	164-3	10.0	8.49	-15	
	Cal 2	I6-0746	164-4	30.0	27.4	-8.8	
		I6-0775	164-5	30.0	27.3	-9.0	
		I6-0810	164-6	30.0	27.2	-9.2	
	Cal 3	I6-0747	164-7	100	104	4.4	
		I6-0776	164-8	100	108	7.6	
		I6-0811	164-9	100	101	0.59	
	Cal 4	I6-0748	164-10	300	315	4.9	
		I6-0777	164-11	300	325	8.5	
		I6-0812	164-12	300	294	-2.0	
	Cal 5	I6-0749	164-13	1000	974	-2.6	
		I6-0778	164-14	1000	1022	2.2	
		I6-0813	164-15	1000	961	-3.9	

Table 15: [ | Calibration Samples - Routine Analyses

Table 15: [ ] Calibration Samples - Routine Analyses							
<u>Sequence</u>	<u>QC</u>	<u>Run #</u>	<u>Ref#</u>	Target Conc	Analyzed Conc	<u>% RE</u>	
		(534006-)	(534006-)	(ng/mL)	(ng/mL)		
I6-534006j	Cal 1	I6-1034	192-1	10.0	8.67	-13	
serum		I6-1072	192-2	10.0	10.4	4.5	
		I6-1117	192-3	10.0	10.5	5.0	
•	Cal 2	I6-1035	192-4	30.0	26.9	-10	
		I6-1073	192-5	30.0	33.7	12	
		I6-1118	192-6	30.0	33.3	11	
	Cal 3	I6-1036	192-7	100	96.2	-3.8	
		I6-1074	192-8	100	99.3	-0.67	
		I6-1119	192-9	100	103	3.2	
	Cal 4	I6-1037	192-10	300	270	-10	
		I6-1075	192-11	300	298	-0.53	
•		I6-1120	192-12	300	302	0.70	
	Cal 5	I6-1038	192-13	500	465	-7.1	
		I6-1076	192-14	500	499	-0.11	
		I6-1121	192-15	500	517	3.4	
	Cal 6	I6-1039	192-16	750	714	-4.9	
•		I6-1077	192-17	750	763	1.8	
		I6-1122	192-18	750	794	5.9	
	Cal 7	I6-1040	192-19	1000	963	-3.7	
		I6-1078	192-20	1000	1005	0.49	
		I6-1123	192-21	1000	1069	6.9	

# PHARMACOKINETIC (IN BLOOD) AND EXCRETION STUDY OF [ ] IN RATS Table 15: [ ] Calibration Samples - Routing Angle

Table 15: [ ] Calibration Samples - Routine Analyses							
<u>Sequence</u>	<u>QC</u>	<u>Run #</u>	Ref#	Target Conc	Analyzed Conc	<u>% RE</u>	
		(534006-)	(534006-)	(ng/mL)	(ng/mL)		
	•						
I6-534006k	Cal 1	I6-1140	205-1	10.0	9.64	-3.6	
serum		I6-1162	205-2	10.0	8.91	-11	
		I6-1213	205-3	10.0	11.4	14	
	Cal 2	I6-1141	205-4	30.0	29.9	-0.28	
	Ou1 <b>2</b>	I6-1163	205-5	30.0	27.8	-7.4	
		I6-1214	205-6	30.0	32.9	10	
	Cal 3	I6-1142	205-7	100	99.0	-1.0	
		I6-1164	205-8	100	98.6	-1.4	
		I6-1215	205-9	100	107	7.2	
•	Cal 4	I6-1143	205-10	300	287	-4.4	
		I6-1165	205-11	300	279	-7.1	
		I6-1216	205-12	300	313	4.4	
	Cal 5	I6-1144	205-13	500	488	-2.3	
	Our 5	I6-1166	205-14	500	481	-3.7	
		I6-1217	205-15	500	530	6.1	
	Cal 6	I6-1145	205-16	750	720	-4.0	
		I6-1167	205-17	750	718	-4.2	
		I6-1218	205-18	750	795	6.0	
	Cal 7	I6-1146	205-19	1000	981	-1.9	
	Cui /	I6-1140	205-19	1000	971	-2.9	
		I6-1219	205-20	1000	1087	-2.9 8.7	
		10 1217	202-21	1000	1007	0.7	

Table 15: [ ] Calibration Samples - Routine Analyses									
Sequence	<u>QC</u>	Run#	Ref#	Target Conc	Analyzed Conc	<u>% RE</u>			
		(534006-)	(534006-)	(ng/mL)	(ng/mL)				
16-53400611	Cal 1	I6-1270	216-1	10.0	10.6	5.8			
urine		I6-1306	216-2	10.0	11.4	14			
		I6-1346	216-3	10.0	8.08	-19			
	Cal 2	I6-1271	216-4	30.0	28.5	-5.1			
		I6-1307	216-5	30.0	29.2	-2.5			
		I6-1347	216-6	30.0	31.5	4.8			
	Cal 3	I6-1272	216-7	100	101	0.87			
		I6-1308	216-8	100	103	3.1			
		I6-1348	216-9	100	99.2	-0.82			
·	Cal 4	I6-1273	216-10	300	321	7.1			
		I6-1309	216-11	300	307	2.2			
		I6-1349	216-12	300	300	-0.15			
	Cal 5	I6-1274	216-13	500	511	2.1			
		I6-1310	216-14	500	472	-5.5			
		I6-1350	216-15	500	459	-8.1			
	Cal 6	I6-1275	216-16	750	732	-2.4			
		I6-1311	216-17	750	753	0.46			
		I6-1351	216-18	750	693	-7.7			
	Cal 7	I6-1276	216-19	1000	1132	13			
		I6-1312	216-20	1000	1014	1.4			
		I6-1352	216-21	1000	984	-1.6			

	<u>AN</u>	ID EXCRETION	<u>ON STUDY</u>	OF[]IN RA	TS	
•	Table 1	5: [ ] Calibr	ation Sampl	les - Routine A	nalyses	
<u>Sequence</u>	<u>QC</u>	<u>Run #</u>	<u>Ref#</u>	Target Conc	Analyzed Conc	<u>% RE</u>
		(534006-)	(534006-)	(ng/mL)	(ng/mL)	
I6-534006m	Cal 1	I6-1370	227-1	10.0	9.29	-7.1
urine		I6-1398	227-2	10.0	9.56	-4.4
		I6-1435	227-3	10.0	11.5	15
	Cal 2	*I6-1371	*227-4	30.0	*67.9	*130
		I6-1399	227-5	30.0	28.8	-3.9
		I6-1436	227-6	30.0	27.6	-8.1
	Cal 3	I6-1372	227-7	100	87.8	-12
		I6-1400	227-8	100	106	5.6
		I6-1437	227-9	100	116	16
	Cal 4	I6-1373	227-10	300	312	4.0
		I6-1401	227-11	300	295	-1.8
		I6-1438	227-12	300	291	-3.2
	Cal 5	I6-1374	227-13	500	516	3.3
•		I6-1402	227-14	500	536	7.1
		I6-1439	227-15	500	479	-4.1
	Cal 6	I6-1375	227-16	750	757	0.99
		I6-1403	227-17	750	673	-10
		I6-1440	227-18	750	764	1.8
	Cal 7	I6-1376	227-19	1000	919	-8.1
		I6-1404	227-20	1000	1081	8.1
		I6-1441	227-21	1000	1027	2.7

^{*} Sample not included in regression analysis based on test for outliers.

Table 15: [ ] Calibration Samples - Routine Analyses								
Sequence	<u>QC</u>	<u>Run #</u>	Ref#	Target Conc	Analyzed Conc	<u>% RE</u>		
		(534006-)	(534006-)	(ng/mL)	(ng/mL)			
I6-534006o	Cal 1	I6-1530	250-1	10.0	9.10	-9.0		
urine		I6-1531	250-2	10.0	10.3	3.5		
		I6-1558	250-3	10.0	10.6	6.0		
	Cal 2	I6-1532	250-4	30.0	31.2	4.0		
		I6-1533	250-5	30.0	27.9	-6.8		
		I6-1559	250-6	30.0	31.0	3.2		
	Cal 3	I6-1534	250-7	100	89.4	-11		
		I6-1535	250-8	100	88.6	-11		
		I6-1560	250-9	100	113	13		
	Cal 4	I6-1536	250-10	300	279	-7.2		
		I6-1561	250-11	300	334	11		
		I6-1562	250-12	300	310	3.2		
	Cal 5	I6-1537	250-13	500	496	-0.78		
		I6-1563	250-14	500	529	5.8		
•.		I6-1564	250-15	500	518	3.6		
	Cal 6	I6-1538	250-16	750	703	-6.2		
		I6-1565	250-17	750	754	0.49		
		I6-1566	250-18	750	745	-0.68		
	Cal 7	I6-1539	250-19	1000	960	-4.0		
		I6-1567	250-20	1000	1007	0.74		
		I6-1568	250-21	1000	1022	2.2		

Table 16: [ ] Quality Control Samples - Routine Analyses Sequence <u>QC</u> Run# Ref# Target Conc Analyzed Conc % RE (534006-) (534006-) (ng/mL) (ng/mL) I6-534006f1 QC1 I6-0684 155-2 30.0 28.0 -6.6 16-0723 155-3 30.0 27.4 serum -8.8 QC2 16-0685 155-5 250 217 -13 I6-0724 155-6 250 225 -9.9 QC3 I6-0686 155-8 750 823 9.7 I6-0725 155-9 750 789 5.2 QC4 I6-0687 10000 155-11 9570 -4.3 I6-0726 155-12 10000 8753 -12 I6-534006g QC1 30.0 I6-0751 165-1 27.4 -8.7 urine I6-0780 165-2 30.0 25.8 -14 I6-0804 165-3 30.0 -5.7 28.3 QC2 I6-0752 165-4 250 233 -7.0 I6-0781 165-5 250 206 -18 16-0805 165-6 250 217 -13 QC3 I6-0753 697 165-7 750 -7.1 I6-0782 165-8 750 775 3.4 I6-0806 165-9 750 755 0.69 QC4 I6-0754 165-10 10000 9516 -4.8 I6-0783 165-11 10000 9624 -3.8 I6-0807 165-12 10000 9657 -3.4

	Table 16: [ ] Quality Control Samples - Routine Analyses								
Sequence	QC	Run#	Ref#	Target Conc	Analyzed Conc	% RE			
-		(534006-)	(534006-)	(ng/mL)	(ng/mL)				
	**								
I6-534006j	QC1	I6-1042	193-1	30.0	28.3	-5.6			
serum		I6-1080	193-2	30.0	30.5	1.8			
		I6-1112	193-3	30.0	30.1	0.41			
	QC2	I6-1043	193-4	250	236	-5.8			
		I6-1081	193-5	250	243	-2.9			
		I6-1113	193-6	250	249	-0.45			
	QC3	I6-1044	193-7	750	732	-2.4			
	•	I6-1082	193-8	750	735	-1.9			
		I6-1114	193-9	750	743	-0.94			
	QC4	I6-1045	193-10	30000	27980	-6.7			
		I6-1083	193-11	30000	30449	1.5			
		I6-1115	193-12	30000	30268	0.89			
I6-534006k	QC1	I6-1148	206-1	30.0	30.8	2.5			
serum		I6-1170	206-2	30.0	30.6	1.9			
		I6-1208	206-3	30.0	32.4	8.2			
	QC2	I6-1149	206-4	250	246	-1.7			
	`	I6-1171	206-5	250	242	-3.1			
		I6-1209	206-6	250	261	4.3			
	QC3	I6-1150	206-7	750	734	-2.1			
	~	I6-1172	206-8	750	747	-0.42			
		I6-1210	206-9	750	792	5.5			
	QC4	I6-1151	206-10	30000	29723	-0.92			
	ζυ,	I6-1173	206-10	30000	28014	-6.6			
		I6-11/3	206-11	30000	33381	11			
		<b></b> -	<b></b>						

Table 16: [ ] Quality Control Samples - Routine Analyses									
Sequence	<u>QC</u>	Run#	<u>Ref#</u>	Target Conc	Analyzed Conc	% RE			
		(534006-)	(534006-)	(ng/mL)	(ng/mL)				
16-53400611	QC1	I6-1278	217-1	30.0	27.9	-7.1			
urine	`	I6-1314	217-2	30.0	26.9	-10			
		I6-1336	217-3	30.0	24.9	-17			
	QC2	I6-1279	217-4	250	227	-9.2			
		I6-1315	217-5	250	208	-17			
		16-1337	217-6	250	229	-8.6			
	QC3	I6-1280	217-7	750	770	2.6			
		I6-1316	217-8	750	758	1.1			
		I6-1338	217-9	750	746	-0.54			
	QC4	I6-1281	217-10	30000	28222	-5.9			
	•	I6-1317	217-11	30000	25817	-14			
	·	I6-1339	217-12	30000	26170	-13			
cage rinse	QC1	I6-1283	219-1	30.0	31.2	3.9			
		I6-1319	219-2	30.0	29.4	-2.0			
		I6-1341	219-3	30.0	29.9	-0.24			
	QC2	I6-1284	219-4	250	240	-3.9			
		I6-1320	219-5	250	232	-7.1			
		I6-1342	219-6	250	253	1.2			
	QC3	I6-1285	219-7	750	750	0.062			
	-	I6-1321	219-8	750	783	4.4			
		I6-1343	219-9	750	783	4.4			
	QC4	I6-1286	219-10	30000	27523	-8.3			
		I6-1322	219-11	30000	28519	-4.9			
		I6-1344	219-12	30000	27956	-6.8			

	Table 16: [ ] Quality Control Samples - Routine Analyses								
<u>Sequence</u>	<u>QC</u>	<u>Run #</u>	<u>Ref#</u>	Target Conc	Analyzed Conc	<u>% RE</u>			
		(534006-)	(534006-)	(ng/mL)	(ng/mL)	-			
I6-534006m	QC1	I6-1378	228-1	30.0	28.8	-3.9			
urine		I6-1406	228-2	30.0	28.3	-5.8			
		I6-1430	228-3	30.0	27.5	-8.3			
	QC2	I6-1379	228-4	250	225	-10			
	202	I6-1407	228-5	250	207	-17			
		I6-1431	228-6	250	265	6.1			
	QC3	I6-1380	228-7	750	645	-14			
		I6-1408	228-8	750	683	-8.9			
		I6-1432	228-9	750	818	9.1			
	QC4	I6-1381	228-10	150000	173620	16			
	QU,	I6-1409	228-10	150000	164740	9.8			
		I6-1433	228-12	150000	160180	6.8			
						0.0			
I6-534006o	QC1	I6-1541	251-1	30.0	30.1	0.20			
urine		I6-1542	251-2	30.0	26.9	-10			
	·	I6-1551	251-3	30.0	27.3	-9.0			
	0.00	TC 1540	051.4	250	0.50				
*	QC2	I6-1543	251-4	250	253	1.1			
		I6-1544 I6-1552	251-5 251-6	250	258	3.3			
		10-1332	231-6	250	224	-10			
	QC3	I6-1545	251-7	750	807	7.6			
		I6-1553	251-8	750	736	-1.9			
		I6-1554	251-9	750	732	-2.4			
	QC4	I6-1546	251-10	30000	29442	-1.9			
	ζ	I6-1555	251-11	30000	28811	-4.0			
		I6-1556	251-12	30000	32312	7.7			

#### **FIGURES 1 - 57**

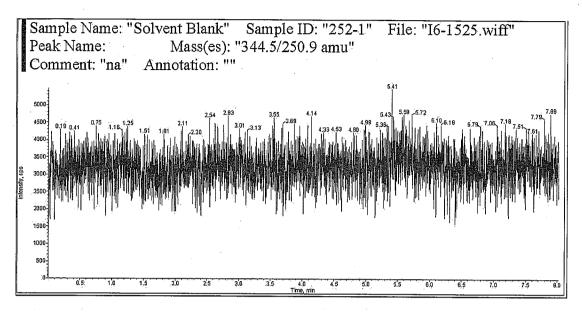


Figure 1: Representative Chromatogram Of A Solvent Blank

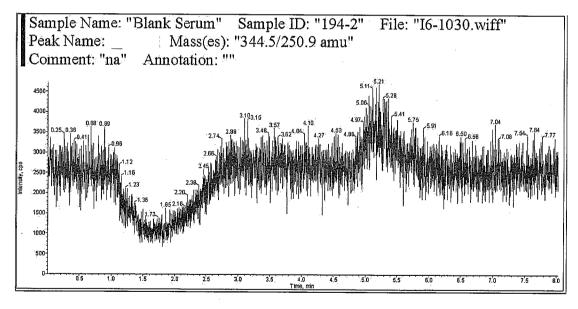


Figure 2: Representative Chromatogram Of A Processed Blank Rat Serum Sample

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## ATTACHMENT I

Supporting Data

Table A-1: 18-534006a Data

Study Record Page: 82b

-87-

Note: Validation session 1

183 of 281

Page 1 of 4

Results Path: \Lcmsp03\sciexdata\Projects\534006\Bio\Results\I8-534006a1.r	
Results Name: I8-534006a1.rdb	

	File Name	Sample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record	Analyte Concentration (ng/mL)	Dilution Factor	Calculated Concentration (ng/mL)
1	l8-534006a\l8-0001.wiff	72-7	Sys Suit	Unknown	•		N/A	1.00	No Intercept
2:	18-534006a\l8-0002.wiff	72-7	Sys Suit	Unknown	***************************************		N/A	1.00	No Intercept
3	I8-534006a\I8-0003.wiff	72-7	Sys Suit	Unknown	••••••		N/A	1.00	No Intercept
4	18-534006a\l8-0004.wiff	72-7	Sys Suit	Unknown			N/A	1.00	No Intercept
5	18-534006a\l8-0005.wiff	71-1	Diluent	Unknown		······	N/A	1.00	No Peak
6	18-534006a\l8-0006.wiff	76-1	Solvent Blank	Unknown		***************************************	N/A	1.00	No Peak
7	18-534006a\l8-0007.wiff	76-2	Serum Blank	Unknown			N/A	1.00	No Peak
8	18-534006a\l8-0008.wiff	76-3	Serum Blank	Unknown			N/A	1.00	2.0660
9	l8-534006a\l8-0009.wiff	76-4	Serum Blank	Unknown			N/A	1.00	No Peak
10	l8-534006a\l8-0010.wiff	71-1	Diluent	Unknown			N/A	1.00	No Peak
11	l8-534006a\l8-0011.wiff	74-1	C 10	Standard		$\boxtimes$	10.000	1.00	10.429
12	I8-534006a\I8-0012.wiff	74-2	C 10	Standard	· ·		10.000	1.00	10.445
13	l8-534006a\l8-0013.wiff	74-3	C 10	Standard		$\boxtimes$	10.000	1.00	9.4050
14	I8-534006a\I8-0014.wiff	74-4	C 30	Standard	<del></del>		30.000	1.00	28.505
15	l8-534006a\l8-0015.wiff	74-5	C 30	Standard		$\boxtimes$	30.000	1.00	28.841
16	18-534006a\l8-0016.wiff	74-6	C 30	Standard	<u> </u>		30.000	1.00	29.016
17	l8-534006a\l8-0017.wiff	74-7	C 100	Standard	ĺ	$\boxtimes$	100.00	1.00	103.90
18	18-534006a\l8-0018.wiff	74-8	C 100	Standard		$\boxtimes$	100.00	1.00	105.07
19	I8-534006a\l8-0019.wiff	74-9	C 100	Standard		$\boxtimes$	100.00	1.00	105.35
20	18-534006a\18-0020.wiff	74-10	C 300	Standard			300.00	1.00	300.75
21	I8-534006a\I8-0021.wiff	74-11	C 300	Standard		×	300.00	1.00	296.48
22	18-534006a\18-0022.wiff	74-12	C 300	Standard	***************************************	$\boxtimes$	300.00	1.00	285.82
23	I8-534006a\I8-0023.wiff	74-13	C 1000	Standard		$\boxtimes$	1000.0	1.00	1015.0
24	I8-534006a\l8-0024,wiff	74-14	C 1000	Standard	**************************************	$\boxtimes$	1000.0	1.00	989.69
25	18-534006a\l8-0025.wiff	74-15	C 1000	Standard	<del></del>	$\boxtimes$	1000.0	1.00	1003.0
26	I8-534006a\l8-0026.wiff	74-16	C 3000	Standard	<del></del>		3000.0	1.00	No Intercept
27	I8-534006a\I8-0027.wiff	74-17	C 3000	Standard	<u></u>		3000.0	1.00	No Intercept
28	I8-534006a\I8-0028.wiff	74-18	C 3000	Standard	<u></u>		3000.0	1.00	No Intercept
29	I8-534006a\I8-0029.wiff	71-1	Diluent	Unknown	***************************************	······	N/A	1.00	No Peak

Printing Date: Wednesday, April 18, 2007 Printing Time: 8:10:35 AM

Page 2 of 4

Results Path: \Lcmsp03\sciexdata\Projects\534006\Bio\Results\18-534006a1.r Results Name: I8-534006a1.rdb

Results	Name: I8-534006a1.rdb	-									
	File Name	Sample ID	Sample Name	Sample Type	Analyte Pe Name		Use Record	Analyte Concentration (ng/mL)	Dilution Factor	Calculated Concentration (ng/mL)	
30	I8-534006a\I8-0030.wiff	75-1	QC 30	Quality Control			Ø	30.000	1.00	28.077	
31	I8-534006a\I8-0031.wiff	75-2	QC 30	Quality Control			$\boxtimes$	30.000	1.00	28.688	
32	I8-534006a\I8-0032.wiff	75-3	QC 30	Quality Control	i		$\boxtimes$	30.000	1.00	29.159	
33	I8-534006a\I8-0033.wiff	75-4	QC 250	Quality Control			$\boxtimes$	250.00	1.00	237.95	
34	I8-534006a\l8-0034.wiff	75-5	QC 250	Quality Control		$\Box$	$\boxtimes$	250.00	1.00	240.98	
35	I8-534006a\I8-0035.wiff	75-6	QC 250	Quality Control	İ		Ø	250.00	1.00	244.27	
36	I8-534006a\I8-0036.wiff	75-7	QC 750	Quality Control			$\boxtimes$	750.00	1.00	713.99	
37	I8-534006a\I8-0037.wiff	75-8	QC 750	Quality Control			$\boxtimes$	750.00	1.00	723.21	
38	I8-534006a\I8-0038.wiff	75-9	QC 750	Quality Control			$\boxtimes$	750.00	1.00	723.44	
39	I8-534006a\I8-0039.wiff	75-10	QC 2500	Quality Control			$\boxtimes$	2500.0	1.00	No Intercept	
40	I8-534006a\I8-0040.wiff	75-11	QC 2500	Quality Control				2500.0	1.00	No Intercept	
41	I8-534006a\I8-0041.wiff	75-12	QC 2500	Quality Control	<u> </u>		***************************************	2500.0	1.00	No Intercept	
42	I8-534006a\I8-0042.wiff	75-13	QC 10000	Quality Control				10000.	20.0	9425.1	
43	I8-534006a\I8-0043.wiff	75-14	QC 10000	Quality Control			$\boxtimes$	10000.	20.0	9383.7	
44	I8-534006a\I8-0044.wiff	75-15	QC 10000	Quality Control		П	$\boxtimes$	10000.	20.0	9490.1	

Printing Date: Wednesday, April 18, 2007 Printing Time: 8:10:35 AM

Page 3 of 4

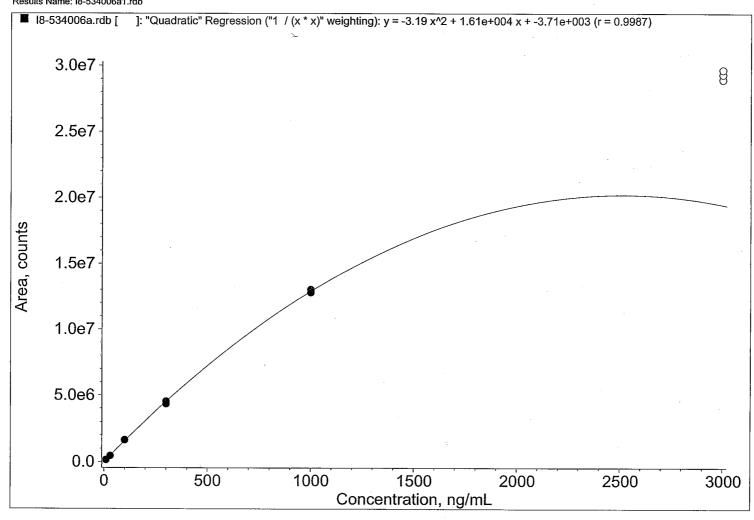
Results Path: \\Lcmsp03\sciexdata\Projects\534006\Bio\Results\l8-534006a1.r Results Name: I8-534006a1.rdb

	File Name	%RE	Analyte Peak Area (counts)	Analyte Retention Time (min)	Analyte Integration Type	Record Modified	Sample Annotation
1	I8-534006a\I8-0001.wiff	#BAD!	3.03e+007	3.98	Base To Base	$\square$	
2	l8-534006a\l8-0002.wiff	#BAD!	3.23e+007	3.71	Base To Base		······································
3	l8-534006a\l8-0003.wiff	#BAD!	3.31e+007	3.66	Base To Base		
4	l8-534006a\l8-0004.wiff	#BAD!	3.34e+007	3.65	Base To Base		
5	I8-534006a\I8-0005.wiff	N/A	0.00e+000	0.00	No Peak		<del></del>
6	18-534006a\l8-0006.wiff	N/A	0.00e+000	0.00	No Peak		,
7	I8-534006a\I8-0007.wiff	N/A	0.00e+000	0.00	No Peak		
8	I8-534006a\I8-0008.wiff	N/A	2.94e+004	3.55	Base To Base		······································
9	I8-534006a\I8-0009.wiff	N/A	0.00e+000	0.00	No Peak		
10	l8-534006a\l8-0010.wiff	N/A	0.00e+000	0.00	No Peak		<del>*************************************</del>
11	l8-534006a\l8-0011.wiff	4.3	1.63e+005	3.52	Base To Base		<del></del>
12	I8-534006a\I8-0012.wiff	4.5	1.64e+005	3.52	Base To Base		······································
13	I8-534006a\I8-0013.wiff	-5.9	1.47e+005	3.51	Base To Base		<del></del>
14	I8-534006a\I8-0014.wiff	-5.0	4.51e+005	3.49	Base To Base		······································
15	I8-534006a\I8-0015.wiff	-3.9	4.57e+005	3.50	Base To Base		·······
16	I8-534006a\I8-0016.wiff	-3.3	4.59e+005	3.50	Base To Base		······································
17	l8-534006a\l8-0017.wiff	3.9	1.63e+006	3.48	Base To Base		······································
18	l8-534006a\l8-0018.wiff	5.1	1.65e+006	3.51	Base To Base		······································
19	I8-534006a\I8-0019.wiff	5.3	1.65e+006	3.46	Base To Base		
20	l8-534006a\l8-0020.wiff	0.25	4.54e+006	3.45	Base To Base		······································
21	l8-534006a\l8-0021.wiff	-1.2	4.48e+006	3.45	Base To Base		
22	l8-534006a\l8-0022.wiff	-4.7	4.32e+006	3.43	Base To Base		
23	l8-534006a\l8-0023.wiff	1.5	1.30e+007	3.43	Base To Base		······································
24	l8-534006a\l8-0024.wiff	-1.0	1.28e+007	3.42	Base To Base		······································
25	l8-534006a\l8-0025.wiff	0.30	1.29e+007	3.41	Base To Base		
26	I8-534006a\I8-0026.wiff	#BAD!	2.89e+007	3.42	Base To Base		
27	I8-534006a\I8-0027.wiff	#BAD!	2.93e+007	3.39	Base To Base		
28	I8-534006a\I8-0028.wiff	#BAD!	2.97e+007	3.38	Base To Base		······································
29	I8-534006a\I8-0029.wiff	N/A	0.00e+000	0.00	No Peak	Tāt	

Printing Date: Wednesday, April 18, 2007 Printing Time: 8:10:35 AM

	File Name	%RE	Analyte Peak Area (counts)		Analyte Integration Type	Record Modified	Sample Annotation
30	I8-534006a\I8-0030.wiff	-6.4	4.45e+005	3.37	Base To Base		
31	I8-534006a\I8-0031.wiff	-4.4	4.54e+005	3.35	Base To Base		······································
32	I8-534006a\I8-0032.wiff	-2.8	4.62e+005	3.36	Base To Base		······································
33	I8-534006a\I8-0033.wiff	-4.8	3.64e+006	3.34	Base To Base		<del></del>
34	I8-534006a\I8-0034.wiff	-3.6	3.68e+006	3.35	Base To Base		***************************************
35	l8-534006a\l8-0035.wiff	-2.3	3.73e+006	3.35	Base To Base		
36	18-534006a\18-0036.wiff	-4.8	9.83e+006	3.36	Base To Base		***************************************
37	I8-534006a\I8-0037.wiff	-3.6	9.94e+006	3.37	Base To Base		······································
38	I8-534006a\I8-0038.wiff	-3.5	9.94e+006	3.35	Base To Base		
39	I8-534006a\I8-0039.wiff	#BAD!	2.44e+007	3.33	Base To Base		······································
40	I8-534006a\I8-0040.wiff	#BAD!	2.40e+007	3.34	Base To Base		······································
41	l8-534006a\l8-0041.wiff	#BAD!	2.46e+007	3.35	Base To Base		
42	8-534006a\ 8-0042.wiff	-5.7	6.85e+006	3.34	Base To Base		······
43	18-534006a\l8-0043.wiff	-6.2	6.83e+006	3.32	Base To Base		······································
44	18-534006a\l8-0044.wiff	-5.1	6.90e+006	3.31	Base To Base		······································

Printing Date: Wednesday, April 18, 2007 Printing Time: 8:10:35 AM



Printing Date: Thursday, March 15, 2007 Printing Time: 10:04:12 AM

Table A-2: I8-534006b Data

Study Record Page: 82c

-93-

Note: Validation session 2

189 of 281

3 Silver	File Name	Sample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record	Analyte Concentration (ng/mL)	Dilution Factor	Calculated Concentration (ng/mL)
1	l8-534006b\l8-0045.wiff	71-1	Diluent	Unknown	•		N/A	1.00	No Peak
2	l8-534006b\l8-0045.wiff	71-1	Diluent	Unknown	••••••		N/A	1.00	No Peak
3	l8-534006b\l8-0045.wiff	71-1	Diluent	Unknown	······	······································	N/A	1.00	No Peak
1	I8-534006b\I8-0046.wiff	81-1	Solvent Blank	Unknown	***************************************		N/A	1.00	< 0
5	I8-534006b\I8-0047.wiff	81-2	Serum Blank	Unknown		***************************************	N/A	1.00	0.17030
	l8-534006b\l8-0048.wiff	81-3	Serum Blank	Unknown			N/A	1.00	0.69883
	l8-534006b\l8-0049.wiff	81-4	Serum Blank	Unknown		······	N/A	1.00	1.0433
	l8-534006b\l8-0050.wiff	71-1	Diluent	Unknown			N/A	1.00	No Peak
	l8-534006b\l8-0051.wiff	79-1	C 10	Standard		×	10.000	1.00	9.9542
0	l8-534006b\l8-0052.wiff	79-2	C 10	Standard		$\boxtimes$	10.000	1.00	10.325
1	l8-534006b\l8-0053.wiff	79-3	C 10	Standard		×	10.000	1.00	9.7624
2	l8-534006b\l8-0054.wiff	79-4	C 30	Standard			30.000	1.00	49.371
3	l8-534006b\l8-0055.wiff	79-5	C 30	Standard	f		30.000	1.00	29.881
4	I8-534006b\I8-0056.wiff	79-6	C 30	Standard	-	$\overline{\boxtimes}$	30.000	1.00	30.016
5	l8-534006b\l8-0057.wiff	79-7	C 100	Standard		$\boxtimes$	100.00	1.00	97.883
6	l8-534006b\l8-0058.wiff	79-8	C 100	Standard		Ø	100.00	1.00	98.277
7	l8-534006b\l8-0059.wiff	79-9	C 100	Standard			100.00	1.00	98.609
8	l8-534006b\l8-0060.wiff	79-10	C 300	Standard		$\boxtimes$	300.00	1.00	317.89
9	l8-534006b\l8-0061.wiff	79-11	C 300	Standard		$\boxtimes$	300.00	1.00	304.62
0	l8-534006b\l8-0062.wiff	79-12	C 300	Standard			300.00	1.00	299.33
1.	I8-534006b\I8-0063.wiff	79-13	C 1000	Standard		× ×	1000.0	1.00	974.30
2	I8-534006b\I8-0064.wiff	79-14	C 1000	Standard	***************************************		1000.0	1.00	982.44
3	I8-534006b\I8-0065.wiff	79-15	C 1000	Standard			1000.0	1.00	1019.5
4	I8-534006b\I8-0066.wiff	79-16	C 3000	Standard			3000.0	1.00	No Intercept
5	l8-534006b\l8-0067.wiff	79-17	C 3000	Standard	[ <del></del>				No Intercept
3	18-534006b\18-0068.wiff	79-18	C 3000	Standard			3000.0	1.00	No Intercept
7	18-534006b\18-0069.wiff	71-1	Diluent	Unknown				1.00	No Peak
3	I8-534006b\l8-0070.wiff	80-1	QC 30	Quality Control			30.000	1.00	31.268
9	I8-534006b\I8-0071.wiff	80-2	QC 30	Quality Control			I		29.209

Printing Date: Wednesday, April 18, 2007 Printing Time: 8:23:11 AM

-95-

	File Name	Sample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record	noitentration	Dilution Factor	Calculated Concentration (ng/mL)
30	I8-534006b\I8-0072.wiff	80-3	QC 30	Quality Control		Ø	30.000	1.00	28.118
31	l8-534006b\l8-0073.wiff	80-4	QC 250	Quality Control	·i	$\boxtimes$	250.00	1.00	238.79
32	l8-534006b\l8-0074.wiff	80-5	QC 250	Quality Control	ı		250.00	1.00	241.12
33	l8-534006b\l8-0075.wiff	80-6	QC 250	Quality Control	***************************************	{	250.00	1.00	247.64
34	I8-534006b\I8-0076.wiff	80-7	QC 750	Quality Control		Ø	750.00	1.00	748.10
35	I8-534006b\I8-0077.wiff	80-8	QC 750	Quality Control		$\boxtimes$	750.00	1.00	753.89
36	I8-534006b\I8-0078.wiff	80-9	QC 750	Quality Control		×	750.00	1.00	746.68
37	I8-534006b\I8-0079.wiff	80-10	QC 2500	Quality Control			2500.0	1.00	No Intercept
38	18-534006b\18-0080.wiff	80-11	QC 2500	Quality Control			2500.0	1.00	No Intercept
39	I8-534006b\I8-0081.wiff	80-12	QC 2500	Quality Control	***************************************		2500.0	1.00	No Intercept
40	I8-534006b\I8-0082.wiff	80-13	QC 10000	Quality Control		$\boxtimes$	10000.	20.0	9708.7
11	I8-534006b\I8-0083.wiff	80-14	QC 10000	Quality Control		$\boxtimes$	10000.	20.0	9684.9
42	I8-534006b\I8-0084.wiff	80-15	QC 10000	Quality Control		$\boxtimes$	10000.	20.0	9625.8

Printing Date: Wednesday, April 18, 2007 Printing Time: 8:23:11 AM

Results Path: \Lcmsp03\sciexdata\Projects\534006\Bio\Results\l8-534006b1.r Results Name: I8-534006b1.rdb

Page 3 of 4

File Name	%RE	Analyte Peak Area (counts)	Analyte Retention Time (min)	Analyte Integration Type	Record Modified	Sample Annotation	
I8-534006b\I8-0045.wiff	N/A	0.00e+000	0.00	No Peak	ПП		
I8-534006b\l8-0045.wiff	N/A	0.00e+000	0.00	No Peak			
I8-534006b\I8-0045.wiff	N/A	0.00e+000	0.00	No Peak			
I8-534006b\I8-0046.wiff	#BAD!	1.27e+003	3.12	Base To Base			
18-534006b\18-0047.wiff	N/A	1.10e+004	3.29	Base To Base		·	
18-534006b\18-0048.wiff	N/A	2.04e+004	3.33	Base To Base			
I8-534006b\I8-0049.wiff	N/A	2.65e+004	3.32	Base To Base			
I8-534006b\I8-0050.wiff	N/A	0.00e+000	0.00	No Peak			
8-534006b\ 8-0051.wiff	-0.46	1.84e+005	3.28	Base To Base		· ·	
I8-534006b\I8-0052.wiff	3.3	1.91e+005	3.30	Base To Base			
I8-534006b\I8-0053.wiff	-2.4	1.81e+005	3.28	Base To Base			
I8-534006b\I8-0054.wiff	65.	8.76e+005	3.28	Base To Base			
I8-534006b\I8-0055.wiff	-0.40	5.36e+005	3.28	Base To Base			
18-534006b\18-0056.wiff	0.053	5.38e+005	3.28	Base To Base		······································	
I8-534006b\l8-0057.wiff	-2.1	1.71e+006	3.27	Base To Base			
I8-534006b\I8-0058.wiff	-1.7	1.72e+006	3.29	Base To Base			
I8-534006b\I8-0059.wiff	-1.4	1.72e+006	3.28	Base To Base			
I8-534006b\l8-0060.wiff	6.0	5.27e+006	3.28	Base To Base			
I8-534006b\I8-0061.wiff	1.5	5.07e+006	3.29	Base To Base		······································	
I8-534006b\I8-0062.wiff	-0.22	4.99e+006	3.27	Base To Base			
8-534006b\ 8-0063.wiff	-2.6	1.37e+007	3.30	Base To Base			
8-534006b\ 8-0064.wiff	-1.8	1.38e+007	3.25	Base To Base		***************************************	
l8-534006b\l8-0065.wiff	1.9	1.42e+007	3.27	Base To Base			
I8-534006b\I8-0066.wiff	#BAD!	3.15e+007	3.27	Base To Base			
l8-534006b\l8-0067.wiff	#BAD!	3.12e+007	3.27	Base To Base		······································	
l8-534006b\l8-0068.wiff	#BAD!	3.17e+007	3.27	Base To Base			
l8-534006b\l8-0069.wiff	N/A	0.00e+000	0.00	No Peak			
l8-534006b\l8-0070.wiff	4.2	5.60e+005	3.27	Base To Base			
I8-534006b\I8-0071.wiff	-2.6	5.24e+005	3.25	Base To Base			

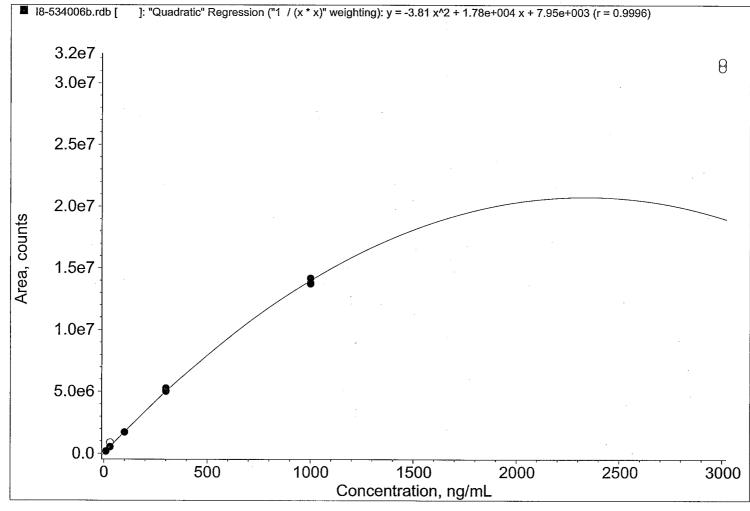
Printing Date: Wednesday, April 18, 2007 Printing Time: 8:23:11 AM

Results Path: \\Lcmsp03\sciexdata\Projects\534006\Bio\Results\\8-534006b1.r Results Name: 18-534006b1.rdb

Page 4 of 4

	File Name	%RE	Analyte Peak Area (counts)		Analyte Integration Type	Record Modified	Sample Annotation
30	I8-534006b\I8-0072.wiff	-6.3	5.05e+005	3.26	Base To Base		
31	I8-534006b\I8-0073.wiff	-4.5	4.04e+006	3.26	Base To Base		
32	I8-534006b\l8-0074.wiff	-3.6	4.07e+006	3.26	Base To Base		······································
33	l8-534006b\l8-0075.wiff	-0.94	4.18e+006	3.25	Base To Base		**************************************
34	I8-534006b\I8-0076.wiff	-0.25	1.12e+007	3.24	Base To Base		······································
35	I8-534006b\I8-0077.wiff	0.52	1.12e+007	3.25	Base To Base		······································
36	I8-534006b\I8-0078.wiff	-0.44	1.12e+007	3.24	Base To Base		<del></del>
37	I8-534006b\I8-0079.wiff	#BAD!	2.70e+007	3.23	Base To Base		
38	I8-534006b\I8-0080.wiff	#BAD!	2.68e+007	3.25	Base To Base		
39	8-534006b\l8-0081.wiff	#BAD!	2.66e+007	3.25	Base To Base		<del></del>
40	l8-534006b\l8-0082.wiff	-2.9	7.74e+006	3.24	Base To Base	lāl	
41	I8-534006b\I8-0083.wiff	-3.2	7.72e+006	3.25	Base To Base		
42	I8-534006b\I8-0084.wiff	-3.7	7.68e+006	3.24	Base To Base		······································

Printing Date: Wednesday, April 18, 2007 Printing Time: 8:23:12 AM



Printing Date: Thursday, March 15, 2007 Printing Time: 10:09:50 AM

194 of 281

-98-

Table A-3: 18-534006g Data

Study Record Page: 129b

Note: Validation session 3

-99-

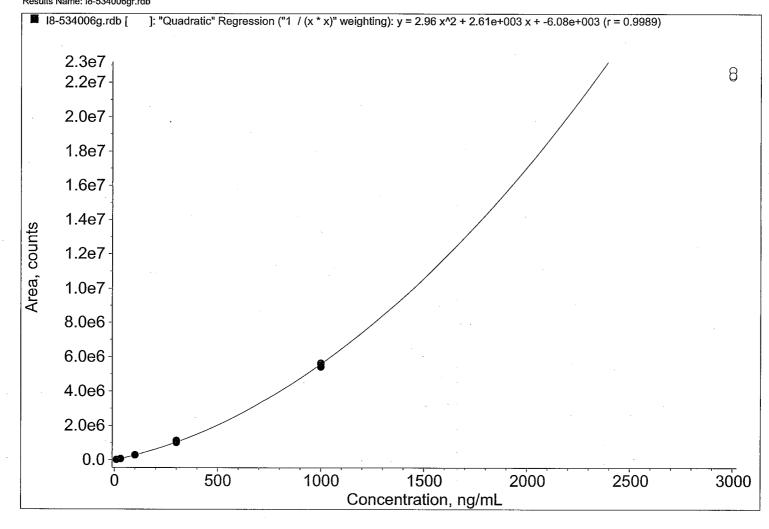
195 of 281

Page 1 of 1

## Results Path: \\Lcmsp03\sciexdata\Projects\534006\Bio\Results\18-534006gr.r Results Name: 18-534006gr.rdb

	File Name	Sample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record	Analyte Concentration (no/mL)	Dilution Factor	Calculated Concentration (po/mL)	%RE	Analyte Peak Area (counts)	Analyte Retention	Analyte Integration Type	Record Modified	Sample Annotation
	18-534006g\18-0257.wiff	115-8	Sys Suit	Unknown	'		N/A	1.00	1995.8	N/A	1.70e+007	Time (min) 3.22	Valley	Ìп	
	18-534006g\18-0258.wiff	115-8	Sys Suit	Unknown	***************************************		N/A	1.00	2027.8	N/A	1.75e+007	3.22	Valley		
	t8-534006g\l8-0259.wiff	115-8	Sys Suit	Unknown	***************************************		N/A	1.00	2853.0	N/A	3.15e+007	3.21	Valley	H	
	18-534006g\18-0260.wiff		Sys Suit	Unknown			N/A	1.00	2837.1	N/A	3.12e+007	3.18	Valley		
******	18-534006g\18-0261,wiff		Sys Suit	Unknown			N/A	1.00	2237.1	N/A	2.07e+007	3.26	Valley	<del> </del>	
	18-534006g\18-0262.wilf		Sys Suit	Unknown	***************************************		N/A	1.00	2141.0	N/A	1.92e+007	3.23	Valley		~~~~
•••••	18-534006g\18-0263.wiff		Sys Suit	Unknown	·		N/A	1.00	3437.8	N/A	4.40e+007	3.28	Base To Base		
			Sys Suit	Unknown			N/A	1.00	2106.8	N/A	1.86e+007	3.15	Valley		
	18-534006q\18-0265.wiff		Sys Suit	Unknown			N/A	1.00	2375.5	N/A	2.29e+007	3.13	Valley		·····
0	18-534006q\18-0266,wiff		Sys Suit	Unknown	***************************************		N/A	1.00	1370.4	N/A	9.13e+006	3.16	i		***************************************
1	18-534006g\l8-0267.wiff		Sys Suit	Unknown			N/A	1.00	1354.9	N/A	8.97e+006	3.15	Base To Base Base To Base		
2	18-534006g\18-0268.wiff		Sys Suit	Unknown	***************************************		N/A	1.00	1357.9	N/A	9.00e+006	3.15	Base To Base		
3	18-534006g\l8-0269.wiff		Sys Suit	Unknown			N/A	1.00	1361.6	N/A			1		
4	I8-534006g\I8-0270.wiff		Diluent	Unknown			N/A	1.00	No Peak			3.17	Base To Base		
5	18-534006g\l8-0271.will		Solvent Blank	in an arman management					L	N/A	0.00e+000	0.00	No Peak		
6	18-534006g\l8-0272,will		Serum Blank	Unknown	***************************************		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
7	18-534006g\18-0272,will 18-534006g\18-0273.wiff			Unknown	***********		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
			Serum Blank	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
8	18-534006g\18-0274.wiff		Serum Blank	Unknown	***************************************		N/A	1.00	No Peak	N/A		0.00	No Peak		
9	18-534006g\18-0275,wiff		Diluent	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
0	18-534006g\\8-0276.will	<u> </u>	C 10	Standard	***************************************	⊠	10.000	1.00	10.100	1.0		3.19	Base To Base		
1	18-534006g\18-0277.wiff		C 10	Standard		⊠		1.00	10.138	1.4		3.12	Base To Base		
2	18-534006g\18-0278.wiff		C 10	Slandard		⊠		1.00	10.395	4.0		3.14	Base To Base		
3	18-534006g\f8-0279.wiff		C 30	Standard	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	⊠	30.000	1.00	27.263	-9.1	6.73e+004	3.13	Base To Base		
4	18-534006g\18-0280.wiff		C 30	Standard	***************************************	⊠	30.000	1.00	29.616	-1.3	7.39e+004	3.12	Base To Base		
5	18-534006g\18-0281.wiff	117-6	C 30	Standard		×	30.000	1.00	26.707	-11.	6.58e+004	3.13	Base To Base		umunuwwww
	18-534006g\18-0282.will	117-7	C 100	Standard		×	100.00	1.00	102.86	2.9	2.94e+005	3.13	Base To Base	П	
7	18-534006g\18-0283.wiff	117-8	C 100	Standard	***************************************	Ø	100.00	1.00	99.069	-0.93	2.82e+005	3,13	Base To Base	ПП	
8	18-534006g\18-0284.wiff	117-9	C 100	Standard		×	100.00	1.00	103.84	3.8	2.97e+005	3.12	Base To Base		
9	18-534006g\t8-0285,wiff	117-10	C 300	Standard	1	×	300.00	1.00	321,99	7.3	1.14e+006	3.14	Base To Base		
0	18-534006gN8-0286.wiff	117-11	C 300	Standard	i	×	300.00	1.00		2.6		3.13	Base To Base		
1 1	18-534006q\18-0287.wiff	117-12	C 300	Standard	i			1.00	293.38	-2.2		3.14	Base To Base		
2	18-534006g\18-0288,wiff	117-13	C 1000	Standard	i	×	1000.0	1.00	1006,7	0.67		3.14	Base To Base		
	18-534006g\18-0289.wiff		C 1000	Standard		×	1000.0	1.00		0.013		3.14	Base To Base		
	18-534006g\18-0290.wiff		C 1000	Standard	i	Ø	1000.0	1.00	980.56			3.13	Base To Base	ᅡ井ᅴ	
	l8-534006g\l8-0291.wilf		C 3000	Standard	i		3000.0	1.00	2365.1			3.12	Base To Base		
	18-534006g\18-0292.wiff		C 3000	Standard			3000.0	1.00	2342.0	-22.		3.14	Base To Base		
	18-534006g\18-0293,wiff		C 3000	Standard				1.00	2345.8	-22.		3.13	Base To Base		
	18-534006g\18-0294.wiff		Diluent	Unknown	i	<u>L.</u>	N/A	1.00		-22. N/A		0.00	No Peak		
9	18-534006g\18-0295.wiff		QC 30	Quality Control	i		30.000	1.00		1,1		3.10			
	18-534006q\18-0296.wiff		QC 30	Quality Control		****	30.000		26,193	-13.		3.12	Base To Base Base To Base	<u> </u>	
	18-534006g\l8-0297.will		QC 30	Quality Control			30.000			-13. 7.1		3.12		<b>.</b>	
	18-534006q\l8-0298.wiff		QC 250	Quality Control				1.00					Base To Base	<u> </u>	
	18-534006g\l8-0299.wiff		QC 250	Quality Control			250.00			7.1		3.12	Base To Base	<u> </u>	
	18-534006g\l8-0300,wiff		QC 250			***************************************		1.00		7.3		3.14	Base To Base		
	18-534006g\18-0301.wiff			Quality Control			250.00					3,12	Base To Base		
			QC 750	Quality Control				1.00		6.4		3,10	Base To Base		
	18-534006g\18-0302.wiff		QC 750	Quality Control			750.00	1.00		3.1		3.12	Base To Base		
	18-534006g\18-0303.wiff		QC 750	Quality Control			750.00	1.00		4.8		3.10	Base To Base		
	18-534006g\18-0304.wiff		QC 2500	Quality Control			2500.0		2148.3	-14.		3.11	Base To Base		
	18-534006g\18-0305.wiff		QC 2500	Quality Control					2160.3	-14.		3.12	Base To Base		
	18-534006g\18-0306.wiff		QC 2500	Quality Control		⊠			2149.1	-14.	1.93e+007	3,12	Base To Base		***************************************
	18-534006g\18-0307.wiff		QC 10000	Quality Control	1	⊠		20.0	17322.	73.	4.48e+006	3.11	Base To Base		
			QC 10000	Quality Control	***************************************	Ø	10000.	20.0	17622.	76.	4.59e+006	3.09	Base To Base		····
3	18-534006g\18-0309.wiff	118-15	QC 10000	Quality Control			10000,	20.0	17381.				Base To Base		·····

Printing Date: Thursday, March 15, 2007 Printing Time: 10:12:34 AM



Printing Date: Thursday, March 15, 2007 Printing Time: 10:13:19 AM

197 of 281

-101-

-102-198 of 281

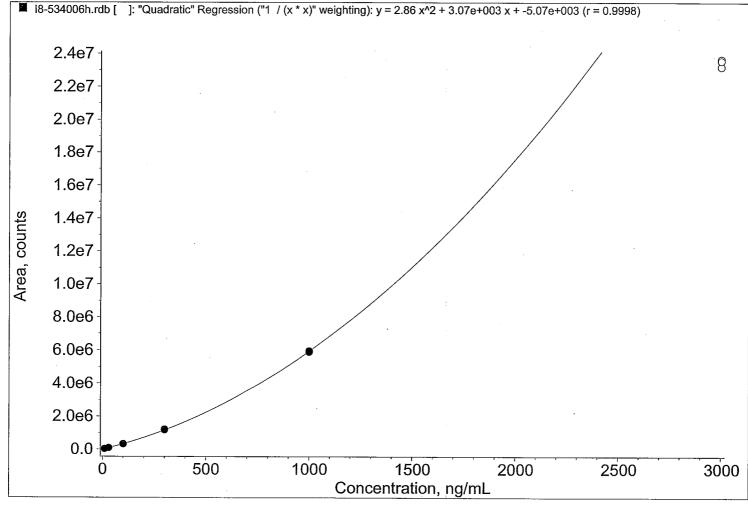
Note:

Validation session 4

Study Record Page: 129c

1	Sample Annotation
2.   8-534006NIB-0311.wiff   124-1   Sohent Blank   Unknown   N/A   1.00   No Peak   N/A   0.00e+000   0.00   No Peak	†
33   86-534006NNB-0312.wiff   124-2   Serum Blank   Unknown   N/A   1.00   No Peak   N/A   0.00e+000   0.00   No Peak	1
S. SAROGINIB-03154.wiff   124-4   Serum Blank   Unknown   N/A   1.00   No Peak   N/A   0.00e+000   0.00   No Peak	1
	1
S.   B. 534006NN6-0317.wif   122-2   C 10   Standard   C   10.000   1.00   10.222   2.2   2.66e+004   3.09   Base To Base   C   10.000   1.00   10.000   1.00   10.000   1.00   1.000   1.00   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000   1.000	
Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Section   Sect	
10	***************************************
11   16   534006NiB-0320.wif   122.5   C 30   Standard   Standa	
11.   B-534006NiB-0320.wif    122-5   C 30   Standard   Standar	1
12.   18-534006NiB-0322.wiff   122-6   C 30   Standard   Standa	
14   18-534006Ni8-0323.wif   122-8   C 100   Standard   S 100.00   1.00   98.208   -1.8   3.24e+005   3.09   Base To Base	<u> </u>
14   18-534006Nile-0323.wif    122-8   C 100   Standard   Standa	
15   8-534006Nik-0324.wif   122-9   C 100   Standard	
16   8-534006Nile 0325.wif    122-10   C 300   Standard	
17.   B-534006NiB-0326.wif    122-11   C 300   Standard	<u> </u>
18   B.534006NiNe 0327.wiff   122-12   C 300   Standard   □ 300.00   1.00   300.26   3.1   1.22e+1006   3.11   Base To Base □	
19   B-534006NiB-0328.wiff   122-13   C 1000   Standard   □ 1000.0   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00	
1000.0   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00   1.00	
21 : IB-534006NIB-0333.wiff   122-15   C 1000   Standard   □ 3000.0   1.00   996.85   0.31   5.90e+006   3.09   Base To Base □ □ 22 : IB-534006NIB-0331.wiff   122-16   C 3000   Standard □ 3000.0   1.00   2381.2   -21   2.32e+007   3.19   Base To Base □ □ 23 : IB-534006NIB-0333.wiff   122-18   C 3000   Standard □ 3000.0   1.00   2383.0   -21   2.32e+007   3.09   Base To Base □ □ 24 : IB-534006NIB-0333.wiff   122-18   C 3000   Standard □ 3000.0   1.00   2383.0   -21   2.36e+007   3.09   Base To Base □ □ 25 : IB-534006NIB-0333.wiff   122-18   C 3000   Standard □ 3000.0   1.00   2383.0   -21   2.36e+007   3.09   Base To Base □ □ 25 : IB-534006NIB-0333.wiff   122-1   C 30   Quality Control □ 30.000   1.00   26.686   -11   7.88e+004   3.10   Base To Base □ □ 27 : IB-534006NIB-0335.wiff   122-1   C 30   Quality Control □ 30.000   1.00   27.454   -8.5   8.13e+004   3.09   Base To Base □ □ 28 : IB-534006NIB-0335.wiff   123-3   C 30   Quality Control □ 30.000   1.00   27.454   -8.5   8.13e+004   3.09   Base To Base □ □ 29 : IB-534006NIB-0335.wiff   123-3   C 30   Quality Control □ 30.000   1.00   27.454   -8.5   8.13e+004   3.12   Base To Base □ □ 30.000   1.00   27.454   -8.5   8.13e+004   3.09   Base To Base □ □ 30.000   1.00   27.454   -8.5   8.39e+004   3.12   Base To Base □ □ 30.000   1.00   27.454   -8.5   8.39e+005   3.00   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000   3.000	
22   8-534006NiB-0333.wif   122-16   0 3000   Standard   3000.0   1.00   2381.2   21.   2.32e+007   3.11   Base To Base   23.   8-534006NiB-0332.wif   122-17   0 3000   Standard   3000.0   1.00   2383.0   24.   2.36e+007   3.09   Base To Base   24.   8-534006NiB-0333.wif   122-18   0 3000   Standard   3000.0   1.00   2383.0   21.   2.36e+007   3.09   Base To Base   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25.   25	
23   18.634006NiB-0332.wiff   122.17   0 3000   0   0   0   0   0   0   0	
24   B.534006Ni8-0333.wiff   122-18   C 3000   Standard   □ 3000.0   1.00   2383.0   21, 2.36e+007   3.09   Base To Base □ □ 25   B-534006Ni8-0334.wiff   114-1   Diluent   Ulrknown   ⊠ 30.000   1.00   2883.0   21, 2.36e+007   3.09   Base To Base □ □ 26   B-534006Ni8-0335.wiff   122-1   CC 30   Quality Control   ⊠ 30.000   1.00   26.686   -11, 7.88e+004   3.10   Base To Base □ □ 27   B-534006Ni8-0335.wiff   122-2   CC 30   Quality Control   ⊠ 30.000   1.00   27.454   8.5   8.13e+004   3.09   Base To Base □ □ 28   B-534006Ni8-0335.wiff   123-3   C 30   Quality Control   ⊠ 30.000   1.00   27.454   8.5   8.13e+004   3.10   Base To Base □ □ 29   B-534006Ni8-0335.wiff   123-3   C 30   Quality Control   ⊠ 30.000   1.00   25.76   2.3   9.67e+005   3.09   Base To Base □ □ 30   B-534006Ni8-0335.wiff   123-5   CC 250   Quality Control   ⊠ 250.00   1.00   255.76   2.3   9.67e+005   3.09   Base To Base □ □ 30   B-534006Ni8-0335.wiff   123-5   CC 250   Quality Control   ⊠ 250.00   1.00   260.38   4.2   9.88e+005   3.10   Base To Base □ □ 30   Base To Base □ 30   Base To Base □ 30   Base To Base □ 30   Base To Base □ 30   Base To Base □ 30   Base To Base □ 30   Base To Base □ 30   Base To Base □ 30   Base To Base □ 30   Base To Base □ 30   Base To Base □ 30   Base To Base □ 30   Base To Base □ 30   Base To Base □ 30   Base To Base □ 30   Base To Base □ 30   Base To Base □ 30   Base To Base □ 30   Base To Base □ 30   Base	***************************************
18-54006NiB-03354.wif   114-1   Dituent   Unknown   N/A   1.00   No Peak   N/A   0.00s+0000   0.00   No Peak	<u> </u>
26   B-534006NiB-0335.wiff   123-1   QC 30   Quality Control   ∑   30.000   1.00   26.686   -11.   7.88e+004   3.10   Base To Base   □   27   B-534006NiB-0335.wiff   123-2   QC 30   Quality Control   ∑   30.000   1.00   27.454   -8.5   8.13e+004   3.09   Base To Base   □   28   B-534006NiB-0337.wiff   123-3   QC 30   Quality Control   ∑   30.000   1.00   25.76   2.3   9.67e+005   3.09   Base To Base   □   29   B-534006NiB-0338.wiff   123-4   QC 250   Quality Control   ∑   250.00   1.00   255.76   2.3   9.67e+005   3.09   Base To Base   □   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.086   30.08	
27   B-534006NiB-0335.wiff   123-2   CC 30   Quality Control   ⊠   30.000   1.00   27.454   8.5   8.13e+004   3.09   Base To Base   □   28   B-534006NiB-0337.wiff   123-3   CC 30   Quality Control   ⊠   30.000   1.00   30.311   1.0   9.05e+004   3.12   Base To Base   □   29   B-534006NiB-0338.wiff   123-4   CC 250   Quality Control   ⊠   250.00   1.00   255.76   2.3   9.07e+005   3.09   Base To Base   □   30.006NiB-0339.wiff   123-5   CC 250   Quality Control   ⊠   250.00   1.00   260.38   4.2   9.88e+005   3.10   Base To Base   □	
28   B-534006NI8-0337.wif   123-3   QC 30   Quality Control   ⊠   30.000   1.00   30.311   1.0   9.05e+004   3.12   Base To Base   □   29   B-534006NI8-0338.wif   123-4   QC 250   Quality Control   ⊠   250.00   1.00   255.76   2.3   9.67e+005   3.09   Base To Base   □   30.000   1.00   260.38   4.2   9.88e+005   3.10   Base To Base   □	
29   B-534006Ni8-0338.wiff   123-4   QC 250   Quality Control   \( \Sqrt{250.00} \)   250.00   1.00   255.76   2.3   9.67e+005   3.09   Base To Base   \( \sqrt{300} \)   8-534006Ni8-0338.wiff   123-5   QC 250   Quality Control   \( \Sqrt{250.00} \)   250.00   1.00   260.38   4.2   9.88e+005   3.10   Base To Base   \( \sqrt{300} \)	·
30 B-534006Nik-0338.wiff 123-5 QC 250 Quality Control 🔯 250.00 1.00 260.38 4.2 9.88e+005 3.10 Base To Base 🔲	
100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 100 CH 10	
31   18-534006h\18-0340.wiff   123-6   QC 250   Quality Control   🖂   250.00   1.00   264.58   5.8   1.01e+006   3.10   Base To Base   🖂	
32   IB-534006hN8-0341.wiff   123-7   QC 750   Quality Control   🔯 750.00   1.00   804.68   7.3   4.32e+006   3.11   Base To Base	<b></b>
33 8-534006hil8-0342.will 123-8 IQC 750 Quality Control 🔯 750.00 1.00 814.52 8.6 4.39e+006 3.10 Base To Base 🖂	
34  B-534006hlk-0343.wiff   123-9   QC 750   Quality Control   🔯 750.00   1.00 798.79   6.5   4.27e+006   3.11   Base To Base	<b> </b>
35  8-534006ini8-0344.wif   123-10  QC 2500  Quality Control   🗵   2500.0   1.00   2199.7   12.   2.06e+007   3.11   Base To Base	ł
36   8-534006hl8-0345.wif   123-11   CC 2500   Quality Control   🖂   2500.0   1.00   2172.5   -13.   2.02e+007   3.09   Base To Base   🗍	İ
37.   16-534006h\ts-0346.wlf   123-12   QC 2500   Quality Control   🔯 2500.0   1.00   2166.0   -13.   2.01e+007   3.10   Base To Base   🗍	
38   8-534006ix 8-0347.wifi   123-13   QC 10000   Quality Control   X   10000.   20.0   11265.   13.   2.63e+006   3.10   Base To Base	
39   8-534006Nl8-0348.wilf   123-14   QC 10000   Quality Control   🔯   10000.   2.0   10601.   6.0   2.43e+006   3.09   Base To Base	ļ
40 I8-534006hil8-0349.wiff   123-15   CC 10000   Quality Control   X   10000.   20.0   10886.   8.9   2.51e+006   3.09   Base TO Base	ļ

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200 of 281

-104-

-105-201 of 281

Note: Validation session 5

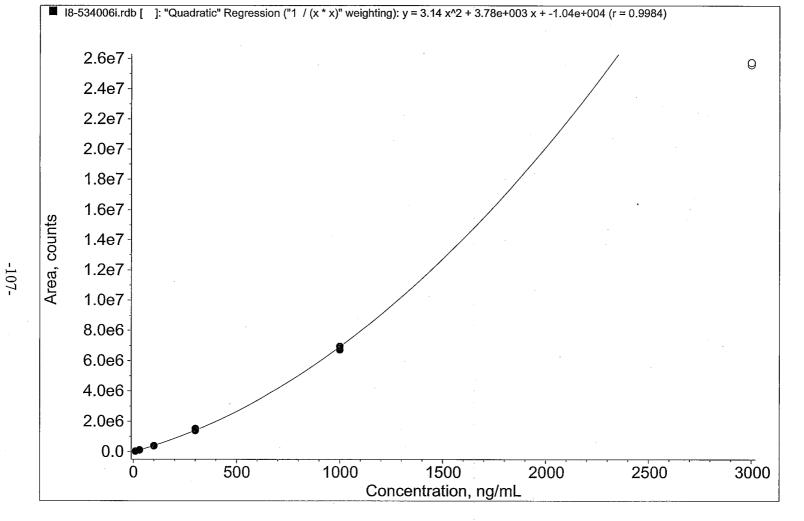
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Page 1 of 1

Results Path: \\Lcmsp03\sciexdata\Projects\534006\Bio\Resuits\18-534006ir.r Results Name: I8-534006ir.rdb

Kes	ults Name: 18-53	40061	.rab												
	File Name	Sample ID	Sample Name	Sample Type	/te Peak ame	Use Record ⁽	Analyte Concentration (ng/ml.)	Dilution Factor	Calculated Concentration (no/mL)	%RE	Analyte Peak Area (counts)	Analyte Retention Time (min)	Analyte Integration Type	Record Modified	Sample Annotation
1	18-534006N18-0350.wiff	114-1	Diluent	Unknown	funitional and	1	N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak	П	
2	18-534006i\18-0351.will	129-1	Solvent Blank	Unknown	1	1	N/A	1.00	3.3146	N/A	2.16e+003	3.17	Base To Base		
3	18-534006i\18-0352.wiff	129-2	Serum Blank	Unknown		1	N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
4	18-534006i\t8-0353.will	129-3	Serum Blank	Unknown		1	N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
5	18-534006i\f8-0354.will	129-4	Serum Blank	Unknown		1	N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
6	18-534006i\18-0355,will	114-1	Diluent	Unknown		1	N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
7	18-534006i\t8-0356.will	127-1	C 10	Standard	i	⊠ 1	10.000	1.00	10.029	0.29	2.78e+004	3.09	Base To Base		***************************************
8	18-534006i\18-0357.will	127-2	C 10	Standard			10.000	1.00	10.752	7.5	3.06e+004	3.10	Base To Base		**************************************
9	18-534006i\18-0358.wilf	127-3	C 10	Standard			10.000	1.00	9.2923	-7.1	2.50e+004	3.11	Base To Base		
10	18-534006N18-0359.wiff	127-4	C 30	Standard			30.000	1.00	28.775	-4.1	1.01e+005	3.09	Base To Base		
11	18-534006N18-0360,wiff	127-5	C 30	Standard			30,000	1.00	29.126	-2.9	1.02e+005	3.08	Base To Base	H	
12	18-534006i\18-0361.wiff	127-6	C 30	Standard			30.000	1.00	32.355	7.8	1.15e+005	3.06	Base To Base		**************************************
13	18-534006i\18-0362.wiff	127-7	C 100	Standard			100.00	1.00	93.989	-6.0		3.08	Base To Base		
14	18-534006718-0363.wiff	127-8	C 100	Standard			100.00	1.00	93.837	-6.2	3.72e+005	3.11	Base To Base		
15	18-534006iN8-0364,wiff	127-9	C 100	Standard			100.00	1.00	98.054	-1.9		3.12	Base To Base		
16	18-534006N18-0365.wiff	127-10	C 300	Standard			300.00	1.00	320.45	6.8		3.10	Base To Base		
17	18-534006N8-0366.wiff	127-11	C 300	Standard			300.00	1.00	321.67	7.2	<u> </u>	3.09	Base To Base		***************************************
18	18-534006i\18-0367.wiff	127-12	C 300	Standard	~~~~		300.00	1.00	294.92	-1.7		3.10	Base To Base		***************************************
19	18-534006i\18-0368,wiff	127-13	C 1000	Standard			1000.0	1.00	1002.6	0.26		3.11	Base To Base		
20	18-534006i\18-0369.wiff	127-14	C 1000	Standard			1000.0	1.00	979.31	-2.1		3.09	Base To Base		
21	18-534006A18-0370.wiff	127-15	C 1000	Standard			1000.0	1.00	1002.8	0.28		3.09	Base To Base		^
22	18-534006i\18-0371.wiff	127-16	C 3000	Standard			3000.0	1.00	2314.3	-23.		3.08	Base To Base		
23	18-534006i\18-0372.wiff	127-17	C 3000	Standard			3000.0	1.00	2324.3	-23.		3.10	Base To Base		
24	18-534006i\18-0373.wiff	127-18	C 3000	Standard	*		3000.0	1.00	2322.6	-23.		3.08	Base To Base	-H-	
25	IB-534006i\i8-0374.will	114-1	Diluent	Unknown			WA.	1.00	No Peak	N/A		0.00	No Peak		
26	18-534006N/8-0375,wiff	128-1	QC 30	Quality Control			30,000	1.00	27.671	-7.8	1	3.07	Base To Base		
27	18-534006N8-0376.wiff	128-2	QC 30	Quality Control			30.000	1.00	25,160	-16.		3.10	Base To Base		
28	18-534006N8-0377.wiff	128-3	QC 30	Quality Control			30.000	1.00	28.886	-3.7	. <del> </del>	3.09	Base To Base		
29	18-534006i\t8-0378.wiff	128-4	QC 250	Quality Control			250.00	1.00	245.08	-2.0		3.10	Base To Base		
30	18-534006i\18-0379.wiff	128-5	QC 250	Quality Control			250,00	1.00	246.92	-1.2		3.08	Base To Base		
31	18-534006i\18-0380.will	128-6	QC 250	Quality Control			250.00	1.00	271.40	8.6		3.10	Base To Base		
32	18-534006N8-0381.wiff	128-7	QC 750	Quality Control	***************************************		750.00	1.00	787.59	5.0		3.10	Base To Base		
33	18-534006i\18-0382.vviff	128-8	QC 750	Quality Control			750.00	1.00	806.53	7.5		3.10	Base To Base	H	
34	18-534006i\18-0383.wiff	128-9	QC 750	Quality Control			750.00	1.00	825.52	10.	£	3.08	Base To Base		
35	18-534006NI8-0384.wiff	128-10	QC 2500	Quality Control			2500.0	1.00	2187.1	-13.	<b></b>	3.09	Base To Base		
36	18-534006i\18-0385.wiff	128-11	QC 2500	Quality Control			2500.0	1.00	2164.3	-13.		3.09	Base To Base		
37	18-534006/\l8-0386.wiff	128-12	QC 2500	Quality Control			2500.0	1.00	2087.2	-17.		3.08	Base To Base		
38	18-534006i\18-0387.wiff	128-13	QC 10000	Quality Control			10000.	20.0	10142.	1,4	i	3.07	Base To Base		
39	18-534006/\l8-0388.wiff	128-14	QC 10000	Quality Control			10000.	20.0	10242.	2.4		3.07	Base To Base		
40	18-53400GN8-0389.wiff		QC 10000	Quality Control			10000,	20.0	10656.	6.6	£	3.07	Base To Base	$\vdash \vdash \vdash \vdash$	
10	III. POPOORIIO OOO WIII	3120-13	140 10000	[scality Collid)		_ ⊠ !'	10000,	20.0	10000,	0.0	2.0051000	J.UI	Person in Days		

Printing Date: Thursday, March 15, 2007 Printing Time: 10:19:09 AM



Printing Date: Thursday, March 15, 2007 Printing Time: 10:19:37 AM

203 of 281

Table A-6: I8-534006j Data

-108-

Note: Validation session 6

204 of 281

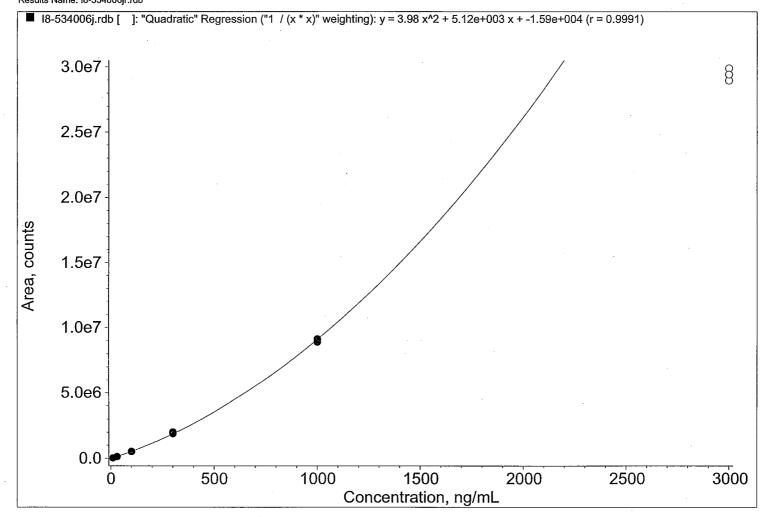
Study Record Page: 142a

Page 1 of 1

	File Name	Sample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record	Analyte Concentration (no/mL)	Dilution Factor	Calculated Concentration (no/mL)	%RE	Analyte Peak Área (counts)	Analyte Retention Time (min)	Analyte Integration Type	Record Modified	Sample Annotation
1	18-534006j\18-0390.wiff	137-8	Sys Suit	Unknown	EEA	Î	N/A	1.00	2012.9	N/A	2.64e+007	3,57	Base To Base		
2	18-534006j\18-0391.wiff	137-8	Sys Suit	Unknown	EEA		N/A	1.00	2020,4	N/A	2.66e+007	3.60	Base To Base		
3	18-534006j\18-0392.wiff	137-8	Sys Suit	Unknown	EEA		N/A	1.00	2038.0	N/A	2.70e+007	3.57	Base To Base		
4	18-534006j\18-0393.wiff	137-8	Sys Suit	Unknown	EEA		N/A	1.00	2039.5	N/A	2.70e+007	3,57	Base To Base		
5	18-534006j\18-0394.wiff	130-1	Diluent		EEA		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
	18-534006j\18-0395.wiff	135-1	Solvent Blank	Unknown	EEA		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
7	18-534006j\18-0396.wiff	135-2	Serum Blank	Unknown	EEA	i	N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
8	18-534006j\18-0397.wiff	135-3	Serum Blank	Unknown	EEA		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
9	18-534006j\18-0398.wiff	135-4	Serum Blank	Unknown	EEA		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
10	18-534006j\18-0399.wiff	130-1	Diluent	Unknown	EEA		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
11	18-534006j\18-0400.wiff		C 10	Standard	EEA	⊠	10.000	1.00	9.9558	-0.44	3.55e+004	3.47	Base To Base		
12 :	18-534006j\18-0401.wiff	133-2	C 10	Standard	EEA	⊠	10.000	1.00	9.4362	-5.6	3.28e+004	3,45	Base To Base		
13	18-534006j\18-0402.wiff	133-3	C 10	Standard	EEA	⊠	10.000	1.00	10.922	9.2	4.05e+004	3.44	Base To Base		
14	18-534006j\18-0403.wiff	133-4	C 30	Standard	EEA	⊠	30.000	1.00	29.732	-0.89	1.40e+005	3.47	Base To Base		
15	18-534006j\18-0404.wiff	133-5	C 30	Standard	EEA	⊠	30.000	1.00	28.407	-5.3	1.33e+005	3.43	Base To Base		
16	18-534006j\i8-0405.wiff	133-6	C 30	Standard	EÉA	×	30.000	1.00	29.197	-2.7	1.37e+005	3.43	Base To Base		
17	18-534006j\18-0406.wiff	133-7	C 100	Standard	EEA	×	100.00	1.00	98.731	-1.3	5.28e+005	3.43	Base To Base	ī ī	
18	18-534006j\18-0407.wiff	133-8	C 100	Standard	EEA		100.00	1.00	98.811	-1.2	5.29e+005	3,40	Base To Base		***************************************
19	18-534006j\18-0408.wiff	133-9	C 100	Standard	EEA	×	100.00	1.00	97.331	-2.7	5.20e+005	3.42	Base To Base		
20	18-534006/\18-0409,wiff	133-10	C 300	Standard	EEA	×	300.00	1.00	311.02	3.7	1.96e+006	3,40	Base To Base		
21	18-534006/\18-0410.wiff	133-11	C 300	Standard	EEA		300.00	1.00	319.83	6.6	2.03e+006	3.42	Base To Base		
22	18-534006j\18-0411.wiff	133-12	C 300	Standard	EEA		300.00	1.00	301.52	0.51	1.89e+006	3.41	Base To Base		
23	18-534006i\18-0412.wiff	133-13	C 1000	Standard	EEA	×	1000.0	1.00	1002.2	0.22	9.12e+006	3.42	Base To Base	Hai	
24	18-534006j\18-0413.wiff	133-14	C 1000	Standard	EEA	Ø	1000.0	1.00	998.26	-0.17	9.06e+006	3.42	Base To Base		
25	18-534006j\18-0414.wiff	133-15	C 1000	Standard	EEA	×	1000.0	1.00	983.97	-1.6	8.88e+006	3.42	Base To Base		
26	18-534006i\18-0415.wiff	133-16	C 3000	Standard	EEA		3000.0	1.00	2171.9	-28.	2.99e+007	3,41	Base To Base		······
27	18-534006j\18-0416.will	133-17	C 3000	Standard	EEA		3000.0	1.00	2130.4	-29.	2.90e+007	3.42	Base To Base		
28	IB-534006i\I8-0417.wiff	133-18	C 3000	Standard	EEA	H	3000.0	1.00	2150.7	-28.	2.94e+007	3.42	Base To Base		
29	18-534006j\(8-0418.will	130-1	Diluent	Unknown	EEA		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
	18-534006i\l8-0419.wiff	134-1	QC 30		EEA	(⊠)	30,000	1.00	28.720	-4.3	1.34e+005	3.39	Base To Base		
Lancinstance of	18-534006j\18-0420.will	134-2	QC 30	Quality Control			30,000	1.00	28.454	-5.2	1.33e+005	3.40	Base To Base		
32	18-534006j\18-0421.will	134-3	QC 30	Quality Control		(X)	30,000	1.00	26.777	-11.	1.24e+005	3.38	Base To Base		·////
33	IB-534006j\l8-0422.wiff	134-4	QC 250	Quality Control	Cauganan muusaa		250.00	1.00	246.29	-1.5	1.49e+006	3.40	Base To Base		
Lawrence	18-534006j\18-0423.wiff	134-5	QC 250	Quality Control			250.00	1.00	245.64	-1.7	1.48e+006	3,40	Base To Base		
Ł	18-534006j\l8-0424.wiff	134-6	QC 250	Quality Control		⊠	250,00	1.00	248,88	-0.45	1.50e+006	3.37	Base To Base		
36	18-534006j\l8-0425.wiff	134-7	QC 750	Quality Control		X X	750.00	1.00	806.92	7.6	6.71e+006	3.39	Base To Base		
37	18-534006i\t8-0426.wiff	134-8	QC 750	Quality Control		<u> </u>	750.00	1.00	771.88	2.9	6.31e+006	3.39	Base To Base		.,
38	18-534006j\18-0427,wilf	134-9	QC 750	Quality Control		X	750.00	1.00	802.28	7.0	6.65e+006	3.37	Base To Base		
39	18-534006j\18-0428.wiff		QC 2500	Quality Control			2500.0	1.00	1974.4	-21.	2.56e+007	3.38	Base To Base	무늬	
Samining	18-534006j\18-0429.wilf		QC 2500 QC 2500	Quality Control		X	2500.0	1.00	2006.0	-21.	2.63e+007	3.36	Base To Base		
	18-534006j\18-0429.will		QC 2500	Quality Control		Ø	2500.0	1.00	1972.7	-20. -21.	2.55e+007	3.36	Base To Base		
			QC 10000	A		Ø		20.0	12734.	1	ž	ž			,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,
42	18-534006j\18-0431.wiff	A management and	·	Quality Control		⊠	10000.			27.	4.86e+006	3.38	Base To Base		
43	18-534006j\l8-0432.wiff		QC 10000	Quality Control		☒	10000.	20.0	12805.	28.	4.89e+006	3.37	Base To Base		
44	18-534006j\18-0433.will	134-15	QC 10000	Quality Control	EEA		10000.	20.0	12728.	27.	4.86e+006	3.36	Base To Base		

Printing Date: Thursday, March 15, 2007 Printing Time: 10:22:30 AM

Results Path: \\Lcmsp03\sciexdata\Projects\534006\Bio\Results\18-534006jr.r Results Name: I8-534006jr.rdb



Printing Date: Thursday, March 15, 2007 Printing Time: 10:22:56 AM

206 of 281

-110-

Table A-7: I8-5340061 Data

Study Record Page: 150a

-111-

Note: Urine cross-validation session

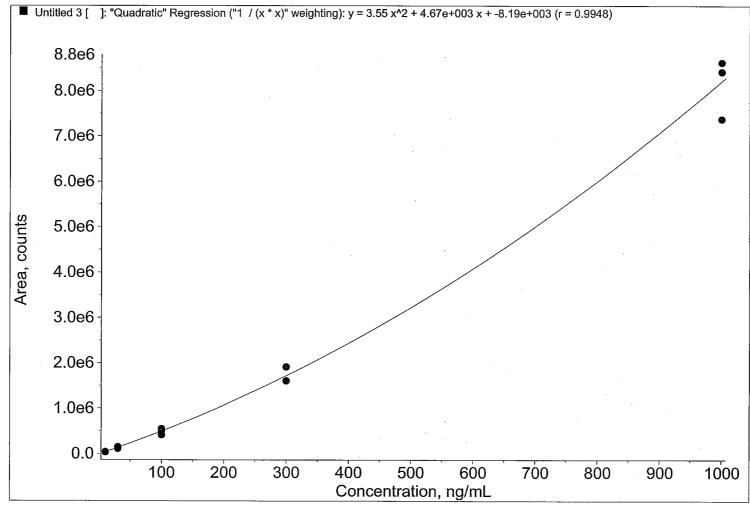
207 of 281

Page 1 of 1

Results Path: \\Lcmsp03\sciexdata\Projects\534006\Bio\Results\18-534006\.rdb Results Name: 18-534006\.rdb

	File Name	Sample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record	Analyte Concentration (ng/mL)	Dilution Factor	Calculated Concentration (ng/mL)	%RE	Analyte Peak Area (counts)	Analyte Retention Time (min)	Analyte Integration	Record Modified	Sample Annotation
1	18-534006N18-0474.wilf	145-8	Sys Suit	Unknown	1	1	N/A	1.00	1803.8	N/A	2.00e+007	3.56	Base To Base	П	
2	18-534006N18-0475.wiff	145-8	Sys Suit	Unknown	1	·	N/A	1.00	2207.8	N/A	2.76e+007	3.37	Base To Base		
3	IB-534006NI8-0476.wiff	145-8	Sys Suit	Unknown	†		N/A	1.00	2212.6	N/A	2.77e+007	3.36	Base To Base		
4	18-534006NI8-0477.wiff	145-8	Sys Suit	Unknown	1		N/A	1.00	2196.8	N/A	2.74e+007	3.35	Base To Base		
5	I8-534006NI8-0478.wiff	143-1	Diluent	Unknown	1		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
6	18-534006N18-0479.wiff	149-1	Solvent Blank	Unknown	Ī	Ĭ	N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
7	18-534006N18-0480.wiff	149-2	Urine Blank	Unknown	1		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
8	18-534006N18-0481.wiff	149-3	Urine Blank	Unknown	1	1	N/A	1.00	2.5302	N/A	3.66e+003	3,39	Base To Base		
9	18-534006N18-0482.wiff	149-4	Urine Blank	Unknown	I ==		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
10.	18-534006N18-0483,wiff	143-1	Diluent	Unknown	1		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
11	18-534006N18-0484.wiff	147-1	C 10	Standard	Ϊ		10.000	1.00	9.3595	-6.4		3.37	Base To Base		
12	18-534006N18-0485.wiff	147-2	C 10	Standard	I	⊠	10.000	1.00	10.563	5.6		3.38	Base To Base		
13	18-534006N18-0486.wiff	147-3	C 10	Standard		⊠	10.000	1.00	10.433	4.3		3.37	Base To Base		
14	18-534006N18-0487.wiff	147-4	C 30	Standard		<u> </u>	30.000		29.185	-2.7		3.37	Base To Base		
15	18-534006N18-0488.wiff		C 30	Standard	I		30.000	1.00	32.801	9.3	1,49e+005	3,39	Base To Base		
16	18-5340061\18-0489.vviff		C 30	Standard		⊠	30.000	1.00	25.068	-16.		3.40	Base To Base		
17	18-534006N18-0490.wiff	.1	C 100	Standard		⋈	100.00	1.00	109.61	9.6		3.35	Base To Base		
18	18-5340061\18-0491.wiff	147-8	C 100	Standard	1	⋈	100.00	1.00	83.967	-16,		3.36	Base To Base		
19	18-5340061\18-0492.wiff		C 100	Standard		⊠	100.00	1.00	98.997	-1.0	.1	3.37	Base To Base		
20	18-534006N18-0493.wiff		C 300	Standard	<u> </u>	⊠	300.00	1.00	328.18	9.4		3,35	Base To Base		
21	18-534006N18-0494.wiff	.4	C 300	Standard			300.00	1.00	283.10	-5.6		3.37	Base To Base		
22	18-5340061\18-0495.wiff		C 300	Standard	L		300.00	1.00	328.25	9.4		3.32	Base To Base		
23	J8-534006I\I8-0496.wiff	1	C 1000	Standard	I	⊠	1000.0	1.00	1034.5	3.5		3.29	Base To Base		
24	18-534006N8-0497,wiff		C 1000	Standard	ļ "	⊠	1000.0	1.00	1017.3	1.7		3.35	Base To Base		
25	18-534006N18-0498.wiff		C 1000	Standard	<u> </u>	⊠	1000.0	1.00	927.04	-7.3		3,35	Base To Base		
26	18-534006N18-0499.wiff		Diluent	Unknown		<b></b>	N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
27	18-534006N8-0500.wiff		QC 30	Quality Control			30.000	1.00	27.883	-7.1	1.25e+005	3.33	Base To Base		
28	18-534006Nt8-0501.wiff		QC 30	Quality Control	ļ		30.000	1.00	36,614	22.	<b></b>	3.37	Base To Base		
29	IB-534006NI8-0502.wiff		QC 30	Quality Control	<u></u>		30.000	1.00	34.717	16.		3.37	Base To Base		
30	18-534006N8-0503.wiff		QC 250	Quality Control	ļ		250.00	1.00	241.83	-3.3		3.37	Base To Base		
31	18-534006N8-0504.wiff		QC 250	Quality Control	ļ	<u></u>	250.00		215.66	-14.		3.37	Base To Base		
32	18-534006N18-0505.will		QC 250	Quality Control	į		250.00	1.00	242.30	-3.1		3.37	Base To Base		
33	18-534006N18-0506.wiff		QC 750	Quality Control	ļ		750.00	1.00		5.7		3.37	Base To Base		
34	18-534006N18-0507.wiff		QC 750	Quality Control	ļ	<u> </u>	750.00	1.00	903.27	20.			Base To Base		
35	18-5340061\18-0508.wiff		QC 750	Quality Control			750.00	1.00		6.8		3.34	Base To Base		
36	18-534006N18-0509.wiff		QC 10000	Quality Control		⊠	10000.	20.0	9977.8	-0.22		3.36	Base To Base		
37	IB-534006NI8-0510.wiff	.3	QC 10000	Quality Control				20.0	12540.	25.		3.33	Base To Base		
38	18-534006N18-0511.wiff	148-12	QC 10000	Quality Control	Lui	⊠	10000.	20.0	10064.	0.64	3,24e+006	3.37	Base To Base		

Printing Date: Friday, March 16, 2007 Printing Time: 7:18:27 AM



Printing Date: Friday, March 16, 2007 Printing Time: 7:19:48 AM

209 of 281

-113-

Note:

Experimental serum sample analysis.

Due to instrument drift only the second half of the run will be reported. The samples from the first half of the run were re-analyzed at a later date.

Table A-8: I6-534006f1 Data

Study Record Page: 161d

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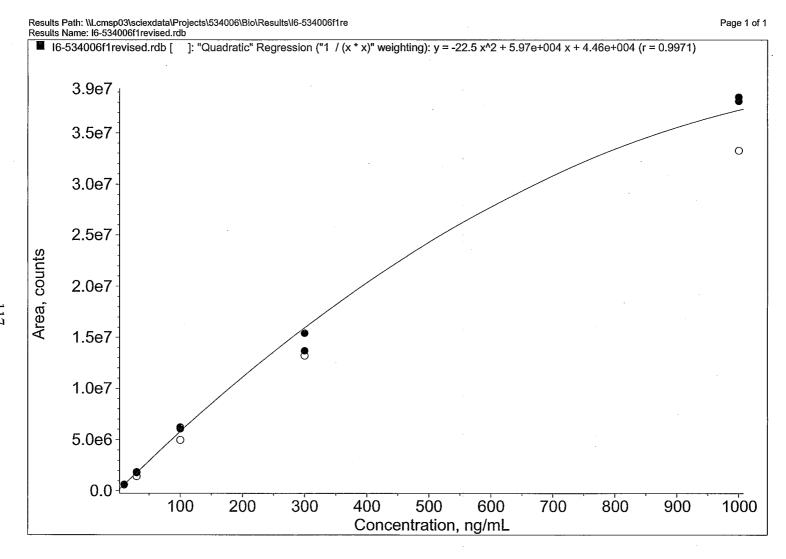
211 of 281

mspos	ovacie)	data\Projects\53	1 34006/BIO/Re	esulis\lb-53	40061	1revis Analyte	1	Calculated	·	T	7	T		I a a second a second a second a second a second a second a second a second a second a second a second a second			F
S	ample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record	Analyte Concentration (ng/mL)	Dilution Factor	Calculated Concentration (ng/mL)	%RE	Analyte Peak Area (counts)	Analyte Retention Time (min)	Analyte Integration Type	Record Modified	Sample Annotation	LCARTRIDGET RAY	LCARTRIDGEP OSITION	
.wiff 15		Fest Solution Fest Solution	Unknown	#****		N/A N/A	1.00	1147,2 980,17	N/A	3.89e+007	5.95	Base To Base	<u> </u>	Managaran da da da da da da da da da da da da da	10	Δ1	
		Test Solution	Unknown	-		N/A		1275.4	N/A N/A	3.69e+007 3.95e+007	5.99 6.01	Base To Base Base To Base			1	B1 C1	
will 15	52-8	lest Solution	Unknown			N/A	1.00	No Intercept	#BAD!	3.96e+007		Base To Base	╁┼			C1	
		Test Solution	Unknown	****		ΝĄ	1.00	1168.1	N/A	3.90e+007	5.95	Valley	┪			Č1	
		lest Solution	Unknown			N/A	1.00	No Intercept	#BAD!	4.03e+007	5.98	Base To Base			1	A1	
		Test Solution Test Solution	Unknown			N/A N/A	1.00	No Intercept No Intercept	#BAD! #BAD!	4.09e+007	6.00 5.93	Base To Base	<u> </u>			B1	
		est Solution	Unknown	;	···········	N/A	1.00	No Intercept	#BAD!	4.10e+007 4.13e+007	5.93	Base To Base Base To Base	<u> </u>			C1	
will 15	52-8	Fest Solution	Unknown			N/A	1.00	No Intercept	#BAD!	4.18e+007		Base To Base	┼┼			C1	
		Mobile Phase	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak	15	***************************************		D1	
		Solvent Blank	Unknown			N/A		0.27302	N/A	6.098+004	5.88	Base To Base				E1	
		Blank Serum Blank Serum	Unknown Unknown	_		N/A N/A	1.00 1.00	0.77502 1.0286	N/A N/A	9,09e+004 1.06e+005	6.01 5.97	Base To Base			J	F1	
		Blank Serum	Unknown			N/A	1.00	1.0162	N/A	1.05e+005	5.94	Base To Base Base To Base				G1 H1	
.wiff 15	51-1 N	Mobile Phase	Unknown	-		N/A		No Peak	N/A	0.00e+000	0.00	No Peak	<del>                                      </del>			G2	
		10	Standard			10.000		9.0798	-9.2	5.85++005	5.94	Base To Base	15		10	C3	
		30	Standard			30.000		23.682	-21.	1.45e+006	5.87	Base To Base				F3	
		300	Standard Standard	·		100.00 300.00		85.103 242.09	-15.	4.96e+006	5.83	Base To Base	<u> </u>			H3	
		31000	Standard	_	뮤	1000.0	1.00	793.60	-19. -21.	1.32e+007 3.32e+007	5.84 5.77	Base To Base Base To Base	<del>  </del>			C4 F4	
		Nobile Phase	Unknown .			N/A		No Peak	N/A	0.00e+000	0.00	No Peak				F4 C6	
	55-1	C 30	Quality Control	****	☒	30.000	1.00	25.040	-17.	1.53e+006	5.75	Base To Base	⊠			E6	
		C 250	Quality Control		☒	250.00		178.95	-28.	1.00e+007	5.72	Base To Base				G6	
		C 750	Quality Control			750.00		571.40	-24.	2.68e+007	5.73	Valley				D7	
		QC 10000 Mobile Phase	Quality Control Unknown		☒	10000. N/A		7066.2 No Peak	-29. N/A	1.83e+007 0.00e+000		Base To Base No Peak				A7	
		6662,D0,1M,TD	Unknown			N/A	1.00	2,5888	N/A	1.99e+005	1	No Peak Base To Base	<u> </u>		11	D1	
		6663,D0,1M,T0	Unknown			N/A	1.00	1.5896	N/A	1.39e+005	5.59	Base To Base	<del></del>			D1	
		6665,D0,1M,T0	Unknown			N/A	1.00	1.7773	N/A	1.51e+005		Base To Base	†===	······································		D1	
		6678,D0,1F,T0	Unknown			N/A		1.3670	N/A	1.26e+005		Base To Base				D1	
		6679,D0,1F,T0	Unknown			WA.		4.7954	N/A	3,30e+005		Base To Base				D1	
		6680,D0,1F,T0 6666.D0.1M.T2 min	Unknown Unknown			N/A N/A	1.00	1.3934 No Intercept	N/A #BAD!	1.28e+005 4.88e+008		Base To Base Base To Base			1:	D1	
		6669,D0,1M,T2 min	Unknown	_		N/A		No Intercept		2.47e+008	10.72	Base To Base				D1	
		6672,D0,1M,T2 min	Unknown	-		N/A		No Intercept		5.27e+008	Ī	Base To Base			I	D1	
		6681,D0,1F,T2 min	Unknown			N/A		No Intercept		5.44e+008		Base To Base			10	D1	
		6684,D0,1F,T2 min	Unknown	4		N/A		No Intercept		5.19e+008		Base To Base		***************************************		D1	
		6685,D0,1F,T2 min 6673,D0,1M,T10min	Unknown Unknown			N/A	1.00	No Intercept No Intercept		5.04e+008 4.72e+008		Base To Base Base To Base		,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	L'	D1	
		6674,D0,1M,T10min	Unknown			WA		No Intercept		4.72e+006 4.56e+008		Base To Base			1	D1 D1	
		6676,D0,1M,T10min	Unknown	-		N/A		No Intercept		4.83e+008		Valley	+H-		1	D1	
		6686,D0,1F,T10min	Unknown			N/A		No Intercept		4.33e+008		Base To Base				D1	
		6688,D0,1F,T10min	Unknown			V/A		No Intercept		3.96e+008		Base To Base				Di	
		6691,DD,1F,T10min 6662,DD,1M,T20 min	Unknown			AVA AVA		No Intercept No Intercept		4.36e+008 4.61e+008		Base To Base			1	D1	
		6663,D0,1M,T20 min	Unknown			VA AV		No Intercept		4.616+008 4.27e+008		Base To Base Base To Base			1	D1	
	7-21 4	6665,D0,1M,T20 min	Unknown			WA		No Intercept		4.55e+008		Valley				51	
			Unknown			ΑVA		No Intercept		3.50e+008		Base To Base				D1	
		6679,D0,1F,T20 min	Unknown	_]		WA AVA		No Intercept		3.31e+008		Base To Base				D1	
will 15		6680,D0,1F,T20 min 6666,D0,1M,T30 min	Unknown			VA VA		No Intercept		3.44e+008		Base To Base				01	
			Unknown			WA AWA	1	No Intercept No Intercept				Base To Base Base To Base	<del>                                     </del>			01	
wiff 150	8-3 4	6672,D0,1M,T30 min	Unknown	-		V/A		No Intercept				Base To Base				01	
		6681,D0,1F,T30 min	Unknown	-		WA AW	1.00	No Intercept	#BAD!	3.42e+008		Base To Base				51	
		6684,D0,1F,T30 min	Unknown			WA		No Intercept			5.44	Base To Base				01	
		6685,D0,1F,T30 min	Unknown			WA.						Base To Base			1.	D1	
		6673,D0,1M,T1 hr 6674,D0,1M,T1 hr	Unknown Unknown					No Intercept				Base To Base				01	
	;		Unknown					No Intercept No Intercept				Base To Base Base To Base				01	
		lobile Phase	Unknown	-+				No Peak				No Peak	- $           -$			D1	
wiff 154	4-2 C	10	Standard	+								Base To Base	무늬			3	
			Standard		X :			29.753									
				工				107.20	7.2								
wiff 154			1	,													
wiff wiff	15	154-5 C 154-8 C 154-11 C	154-5 C 30 154-8 C 100 154-11 C 300	154-5 C 30 Standard 154-8 C 100 Standard 154-11 C 300 Standard	154-5 C 30 Standard 154-8 C 100 Standard 154-11 C 300 Standard	154-5 C 30 Standard ⊠ 3 154-8 C 100 Standard ⊠ 1 154-11 C 300 Standard	154-5 C 30 Standard ⊠ 30,000 154-8 C 100 Standard ☑ 100,00 154-11 C 300 Standard ☑ 300,00	154-5 C 30 Standard ⊠ 30,000 1.00 154-8 C 100 Standard ⊠ 100.00 1.00	154-5         C 30         Standard         S 30.000         1.00         29.753           154-8         C 100         Standard         I 100.00         1.00         107.20           154-11         C 300         Standard         S 300.00         1.00         286.20	154-5         C 30         Standard         ⊠ 30,000         1.00         29,753         -0.82           154-8         C 100         Standard         ⊠ 100,00         1,00         107,20         7,2           154-11         C 300         Standard         ⊠ 300,00         1,00         288,20         -3,9	154-5 C 30 Standard ⊠ 30.000 1.00 29.753 -0.82 1.66e+006 154-8 C 100 Standard ⊠ 100.00 1.00 107.20 7.2 6.19e+006 154-11 C 300 Standard ⊠ 300.00 1.00 286.20 -3.9 1.54e+007	154-5 C 30 Standard ⊠ 30,000 1.00 22,753 -0.82 1.80e+006 5.44 154-8 C 100 Standard ⊠ 100.00 1.00 107.20 7.2 6.19e+006 5.50 154-11 C 300 Standard ⊠ 300.00 1.00 286.20 -3.9 1.54e+007 5.45	154-5 C 30 Standard ⊠ 50.000 1.00 23.753 -0.82 1.80e+006 5.44 Base To Base 154-8 C 100 Standard ⊠ 100.00 1.00 107.20 7.2 6.19e+006 5.50 Base To Base 154-11 C 300 Standard ⊠ 300.00 1.00 288.20 -3.9 1.54e+007 5.45 Base To Base	154-5 C 30 Slandard ⊠ 30.000 1.00 29.753 -0.62 1.60e+006 5.44 6ase To Base □ 154-8 C 100 Slandard ⊠ 100.00 1.00 107.20 7.2 6.19e+006 5.50 6ase □ 154-11 C 300 Slandard ⊠ 300.00 1.00 286.20 -3.9 1.54e+007 5.45 6ase □	154-5 C 30 Standard ☑ 30,000 1.00 29,753 -0.82 1.80e+006 5.44 6ase To Gase ☐ 154-8 C 100 Standard ☑ 100.00 1.00 107.20 7.2 6.19e+006 5.50 Base To Base ☐ 154-11 C 300 Standard ☑ 30,000 1.00 288.20 -3.9 1.54e+007 5.45 Base To Base ☐	154-5 C 30 Standard ⊠ 50,000 1,00 22,753 -0.82 1,80e+006 5,44 Base To Base □ 10 154-8 C 100 Standard ⊠ 100,00 1,00 107,20 7,2 6,19e+006 5,50 Base To Base □ 10 154-11 C 300 Standard ⊠ 300,00 1,00 288.20 -3.9 1,54e+007 5,45 Base To Base □ 10	154-5 C 30 Standard ⊠ 50.000 1.00 22.753 -0.82 1.80e+006 5.44 Base To Base □ 10 F3 154-8 C 100 Standard ⊠ 100.00 1.00 107.20 7.2 6.19e+006 5.50 Base To Base □ 10 B4 154-11 C 300 Standard ⊠ 300.00 1.00 286.20 -3.9 1.54e+007 5.45 Base To Base □ 10 E4

Page 2 of 2

212 of 281

		Sample	<u> </u>	1	Analyte Peak	llee	Analyte	Dilution	Calculated		Analyte Peak	Analyte	Analyte	Record		LCARTRIDGET	LCARTRIDGE
	File Name	aı	Sample Name	Sample Type		Record	Concentration (ng/mL)	Factor	Concentration (ng/mL)	%RE	Area (counts)	Retention Time (min)	Integration Type	Modified	Sample Annotation	RAY	OSITION
7	16-534006N16-0683,will	151-1	Mobile Phase	Unknown	le man		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak	T n		10	C6
8	16-534006A16-0684.wiff	155-2	QC 30	Quality Control		(X)	30,000	1.00	28.021	-6.6	1.70e+006	5.42	Base To Base	n		10	E6
9	16-534006N16-0685.wiff	155-5	QC 250	Quality Control		×	250,00	1.00	216,53	-13.	1.19e+007	5.39	Base To Base			10	G6
0	16-534006N16-0686,wilf	155-8	QC 750	Quality Control		⊠	750.00	1.00	822.58	9.7	3.39e+007	5.36	Base To Base	1 7		10	D7
4	16-534006N16-0687.will	155-11	QC 10000	Quality Control	*****	×	10000.	20.0	9570.4	-4.3	2.35e+007	5.37	Base To Base	1 📅		10	A7
2	16-534006N6-0688.wilf	151-1	Mobile Phase	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak	1 -		10	D1
3	16-534006N6-0689.wiff	158-10	46686,D0,1F,T1 hr	Unknown	*****		N/A	1.00	No intercept	#BAD!	1.45e+008	5.33	Valley	T 7		10	D1
4	16-534006N16-0690,wiff	158-11	46688,D0,1F,T1 hr	Unknown	*****		N/A	1.00	No Intercept	#BAD!	1.97e+008	5.37	Valley	1 7		10	D1
5	16-534006N16-0691,wilf	158-12	46691,D0,1F,T1 hr	Unknown	-		N/A	1.00	No Intercept	#BAD!	2.80e+008	5.30	Base To Base		***************************************	10	D1
6	16-534006N16-0692.wiff	158-13	46662,D0,1M,T3 hr	Unknown	****		N/A	1.00	No Intercept	#BAD!	3.82e+008	5.24	Base To Base	1 7	***************************************	10	D1
7	16-534006N16-0693.wiff	158-14	46663,D0,1M,T3 hr	Unknown			N/A	1.00	No Intercept	#BADI	2.65e+008	5.23	Base To Base	1 7 T		10	D1
8	16-534006N16-0694.wiff	158-15	46665,D0,1M,T3 hr	Unknown	*****		N/A	1.00	No Intercept	#BAD!	3.11e+008	5.21	Base To Base	l H		10	D1
9	16-534006N16-0695.wiff	158-16	46678,D0,1F,T3 hr	Unknown			N/A	1.00	No Intercept	#BAD!	4.65e+007	5.33	Base To Base			10	D1
0	16-534006N16-0696.wiff	158-17	46679,D0,1F,T3 hr	Unknown			N/A	1.00	992.08	N/A	3.71e+007		Base To Base	1-K-	***************************************	10	D1
1	16-534006N16-0697.wiff	158-18	46680,D0,1F,T3 hr	Unknown	******		N/A	1.00	625.06	N/A	2.86e+007		Valley	<u> </u>		10	D1
2	16-534006N16-0698.wiff	158-19	46666,D0,1M,T5 hr	Unknown			N/A	1.00	No Intercept	#BAD!	1.92e+008		Base To Base	<del>                                     </del>	***************************************	10	D1
3	16-534006N16-0699,wiff	158-20	46669.D0,1M,T5 hr	Unknown			N/A	1.00	No Intercept	#BAD!	2.94e+008		Valley	<del>- 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1</del>		10	D1
4	IG-534006NIG-0700.wiff	158-21	46672,D0,1M,T5 hr	Unknown	*****		N/A	1.00	No Intercept	#BAD!	2.78e+008		Base To Base	1-5-		10	D1
5	16-534006N16-0701.wiff	158-22	46681,D0,1F,T5 hr	Unknown			N/A	1.00	638.80	N/A	2.90e+007		Base To Base	1-5-		10	D1
6	16-534006N16-0702.wiff	158-23	46684,D0,1F,T5 hr	Unknown			N/A	1.00	285.27	N/A	1.52e+007		Base To Base	一片一		10	D1
7	6-534006N6-0703,wiff	158-24	46685,D0,1F,75 hr	Unknown		·	N/A	1.00	159.98	N/A	9.02e+006	5.26	Base To Base	<del>                                     </del>		10	D1
8	16-534006N16-0704.wilf	159-1	46673,D0,1M,T7 hr	Unknown			N/A	1.00	No Intercept	#BAD!	3.28e+008		Base To Base	1-H-		10	D1
	16-534006NI6-0705.wiff	159-2	46674,D0,1M,T7 hr	Unknown			N/A	1.00	No Intercept	#BAD!	1.80e+008	5.26	Base To Base	1 #		10	D1
0	6-534006/\(\)16-0706.wiff	159-3	46676,D0,1M,T7 hr	Unknown	•••••		N/A	1.00	No Intercept	#BADI	2.96e+008		Base To Base	1-5-	***************************************	10	D1
		159-4	46686,D0,1F,T7 hr	Unknown	,		N/A	1.00	408.50	N/A	2.07e+007		Base To Base			10	D1
	6-534006N6-0708.wiff	159-5	46688,D0,1F,T7 hr	Unknown			N/A	1.00	77,046	N/A	4.51e+006		Base To Base			10	D1
		159-6	46691,D0,1F,T7 hr	Unknown			N/A	1.67	175.80	N/A	6.09e+006		Base To Base			10	D1
		159-7	46662, D0, 1M, T24 hr	Unknown	·		N/A	1.00	395.41	N/A	2.01e+007		Base To Base	<del>                                     </del>		10	D1
		159-8	46663,D0,1M,T24 hr	Unknown			N/A	1.00	73.163	N/A			Base To Base	1-4-	***************************************	10	D1
	6-534006N6-0712.wiff	159-9	46665,D0,1M,T24 hr	Unknown			N/A	1.00	106.01	N/A	6.12e+006		Base To Base	1-1		10	D1
		159-10	46678,D0,1F,T24 hr	Unknown	·		N/A	1.00	13,162	N/A			Base To Base			10	D1
		159-11	46679,D0,1F,T24 hr	Unknown	,		N/A		9.9031	N/A			Base To Base	1-8-	***************************************	10	D1
	6-534006N6-0715.wiff	159-12	46680,D0,1F,T24 hr	Unknown			N/A	1.00	12.459	NΛ			Base To Base	1 7		10	D1
		159-13	46666,D0,1M,T48 hr	Unknown			N/A	1.00	18.261	NVA			Base To Base			10	D1
		159-14	46669,D0,1M,T48 hr	Unknown			N/A	1.00	178.01	N/A			Base To Base	<del>                                     </del>		10	D1
			46672,D0,1M,T48 hr	Unknown	******		N/A	1.00	22.969	N/A			Base To Base	<del>                                     </del>	······································	10	D1
			46681,D0,1F,T48 hr	Unknown			N/A	1.00	13.444	N/A			Base To Base	<del>                                     </del>		10	D1
			46684,D0,1F,T48 hr	Unknown			N/A	1.00	7.0183	N/A			Base To Base	<del>├</del> ├		10	D1
		159-18	46685,D0,1F,T48 hr	Unknown			N/A	1.00	13.505	N/A			Base To Base	<del>                                     </del>		10	D1
		151-1	Mobile Phase	Unknown			N/A		No Peak	N/A			No Peak	╁╌╫┈		10	C6
		155-3	QC 30	Quality Control		<b>(3)</b>	30.000		27.373	-8.8			Base To Base	l la		10	E6
		155-6	QC 250	Quality Control	•	冈	250.00		225.29	-9.9			Base To Base	N N		10	G6
		155-9	QC 750	Quality Control	•	⊠	750.00	1.00	789.27	5.2			Base To Base	<del></del>		10	D7
			QC 10000	Quality Control		<u> </u>	10000.		8752.6	-12.		L	Base To Base	<del>                                     </del>		10	A7
		151-1	Mobile Phase	Unknown			N/A		No Peak	N/A			No Peak	<del></del>		10	ID1
	6-534006Nt6-0728.wiff	154-3	C 10	Standard			10.000		9.8150	-1.9			Base To Base	<u> </u>	······	10	E3
	6-534006N6-0729.wiff	154-6	C 30	Standard		N N	30,000		31,230	4.1						10	F3
	6-534006N6-0729.will	154-9	C 100	Standard		Ø	I	<u> </u>	104.26				Base To Base				.I
				Standard		×	100.00	1.00		4.3			Base To Base			10	B4
	IIW. I CAD-DINOUDINGGOD	134-12	C 300	:orangarg		×	300.00	1.00	252.08	-16.	1.37e+007	5.26	Base To Base			10	E4



Printing Date: Thursday, March 29, 2007 Printing Time: 6:22:16 PM

-118-214 of 281

Note:

0-6 hour, 6-12 hour and 12-24 hour experimental urine and cage wash samples

Study Record Page: 173a

215	
of 281	117

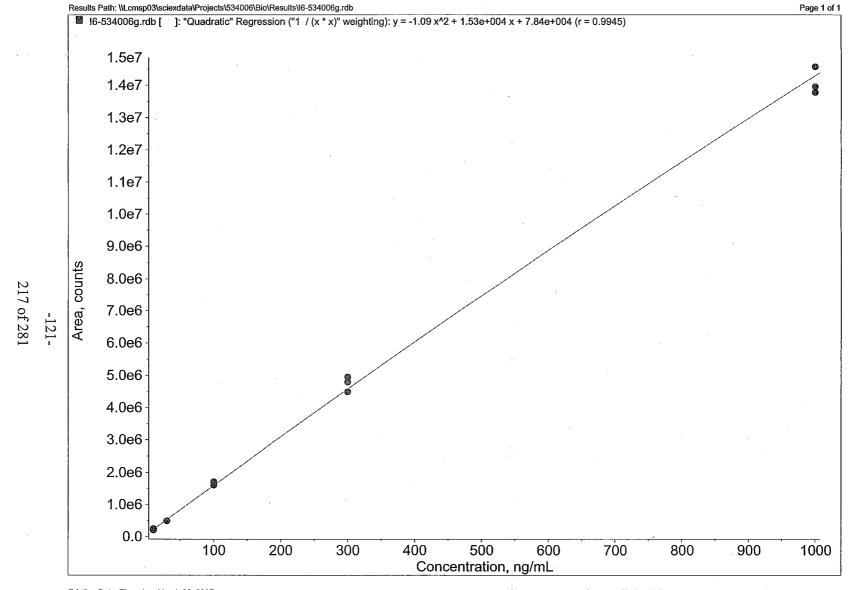
M M	File Name	Sample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record	Analyte Concentration (ng/mL)	Dilution Factor	Calculated Concentration (rig/mL)	%RE	Analyte Peak Area (counts)	Anslyte Retention Time (min)	Analyte Integration Type	Record Modified	Sample Annotation	LCARTRIDGET RAY	LCARTRIDGEP OSITION
7	16-534006g\16-0733.wiff		Test Solution	Unknown	10.	anaman.	NA	1.00	2274.3	N/A	2.93e+007	5.34	Base To Base	To	A PRILITARIA CAL MANAGANA	10	A1
	16-534006g\16-0734,wiff		Test Solution	Unknown				1,00	2199.2	N/A		5.32	Base To Base			10	B1
	16-534006g\16-0735.wiff		Test Solution	Unknown	***			1.00	2234.6	N/A		5,34	Base To Base			10	C1
	16-534006g\16-0736.wiff		Test Solution	Unknown					2251.0	N/A		5.29	Base To Base	<u> </u>		ł	C1
	16-534006g\16-0737.wiff 16-534006q\16-0738.wiff		Test Solution Test Solution	Unknown				1.00 1.00	2210.9 2169.0	N/A N/A		5.37 5.27	Valley Base To Base			10	C1 A1
1	16-534006g\16-0739.wiff		Mobile Phase	Unknown				1.00		N/A		0.00	No Peak				A1 D1
-	16-534006g\16-0740.wiff		Solvent Blank	Unknown				1.00	No Peak	N/A		0.00	No Peak			.t	E1
	16-534006g\16-0741.wiff		Blank Urine	Unknown				1.00	No Peak	NA		0.00	No Peak	├		10	= 1 F1
	16-534006g\16-0742.wiff		Blank Urine	Unknown				1.00	No Peak	NA		0.00	No Peak	1		1	G1
7	16-534006g\16-0743.wiff		Blank Urine	Unknown			N/A	1.00	No Peak	NA	0.00e+000	0.00	No Peak	177		10	HI
~	16-534006g\16-0744.wiff		Mobile Phase	Unknown	***		N/A	1.00	No Peak	N∕A	0.00e+000	0.00	No Peak	l n		10	G2
	16-534006g\16-0745.wiff		C 10	Standard	-	×		1.00		16.		5.43	Base To Base			10	C3
	16-534006g\16-0746.wiff		C 30	Standard					27.360	-8.8		5.54	Base To Base			1	F3
	16-534006g\16-0747.wiff		C 100	Standard				1.00		4.4		5.35	Base To Base			1 1	Н3
_	16-534006g\16-0748.wiff		C 300	Standard						4.9		5.44	Base To Base				C4
	16-534006g\16-0749.wiff 16-534006g\16-0750.wiff		C 1000	Standard		X		1.00	974.28 No Peak	-2.6 N/A		5.43	Base To Base			I	F4
2	16-534006g\16-0751.will		Mobile Phase QC 30	Unknown Quality Control		[C]			No Peak 27.402	-8.7		0.00 5.48	No Peak Base To Base	<u> </u>			C6 E6
4	16-534006g\16-0752.will		QC 250	Quality Control					232.58	-8.7 -7.0		5.47	Base To Base	<u> </u>			⊑6 G6
÷	16-534006gN6-0753.wiff		QC 750	Quality Control					696.98	-7.1		5.49	Base To Base	<del>├-</del> ႘			D7
÷	16-534006g\\6-0754.will		QC 10000	Quality Control					9515.5	-4.8		5.51	Base To Base			11	A7
ँ	16-534006g\16-0755.wilf		Mobile Phase	Unknown	•••		N/A	1.00	No Peak	N/A		0.00	No Peak	<del>-</del> H−		1	D1
Ī	l6-534006g\l6-0756.will	167-1	46664, 0-6 hr	Unknown					No Intercept	#BAD!	5.61e+008	5.41	Base To Base				D1
ं	16-534006g\16-0757.wiff		46664, 0-6 hr rinse	Unknown .			N/A	1.00	No Intercept	#BAD!		5.44	Base To Base			10	D1
Z	16-534006g\16-0758,wiff		46667, 0-6 hr rinse	Unknown			N/A	1.00	No Intercept	#BAD!		5.35	Base To Base				D1
ŝ	16-534006g\16-0759,wiff		46670, 0-6 hr	Unknown					No Intercept	#BAD!		5.52	Valley				D1
	16-534006g\16-0760,will		46670, 0-6 hr rinse	Unknown				1.00	No Intercept	#BAD!		5.40	Base To Base				D1
	16-534006g\16-0761.wiff		46682, 0-6 hr	Unknown			N/A	1.00	No Intercept	#BAD!		5.70	Valley	<u> </u>			D1
	16-534006g\16-0762.wiff 16-534006g\16-0763.wiff		46682, 0-6 hr rinse 46683, 0-6 hr	Unknown Unknown			N/A N/A	1.00 1.00	3948.8 No Intercept	N/A #BAD!		5.46 5.65	Base To Base Base To Base	<u> </u>		F	D1
_	16-534006g\16-0764.wiff		46683, 0-6 hr rinse	Unknown					No Intercept	#BAD!		5,36	Valley	<u> </u>		! · · · · · · · · · · · · · · · · · · ·	D1
÷	16-534006g\16-0765.wiff		46690, 0-6 hr	Unknown					No Intercept	#BAD!		5.47	Valley			I	D1
-	16-534006g\l6-0766.wiff		46690, 0-6 hr rinse	Unknown					No Intercept	#BAD!		5.40	Base To Base			I	D1
			46664, 6-12 hr	Unknown						#BAD!		5,34	Base To Base	<u> </u>			D1
į	16-534006g\16-0768,wiff	167-13	46664, 6-12 hr rinse	Unknown			N/A	1.00	No Intercept	#BAD!	6.26e+007	5.46	Base To Base			10	D1
਼		167-14	46667, 6-12 hr	Unknown			N/A	1.00	No intercept	#BAD!	4.31e+008	5.43	Base To Base			10	D1
7			46667, 6-12 hr rinse	Unknown					4263.3	N/A		5,38	Base To Base			10 (	D1
			46670, 6-12 hr	Unknown				1.00	No Intercept	#BAD!		5.39	Base To Base				D1
			46670, 6-12 hr rinse	Unknown			N/A		3269.5	N/A		5.43	Base To Base				D1
3	16-534006g\l6-0773,wilf 16-534006g\l6-0774.wilf		Mobile Phase C 10	Unknown Standard					No Peak 10.600	N/A 6.0		0.00	No Peak			1	D1
ļ	16-534006g\\6-0775.wilf		C 30	Standard		X			27.309	-9.0		5.56 5.55	Base To Base Base To Base				E3 F3
-			C 100	Standard						7.6		5.57	Base To Base				ra R4
2	16-534006g\\6-0777.wiff		C 300	Standard	-	⊠ ⊠				8.5		5.57	Base To Base		<b></b>	1	E4
			C 1000	Standard		⊠		1.00		2.2		5.51	Base To Base				G4
ä	16-534006g\16-0779.wiff	168a-1	Mobile Phase	Unknown			N/A	1.00		N/A		0.00	No Peak				C6
	16-534006g\16-0780,wiff	165-2	QC 30	Quality Control	•••	⊠	30.000	1.00	25.751	-14.	4.72e+005	5.59	Base To Base			10	E6
Ī			QC 250	Quality Control	•	Ø			206.04	-18.		5.59	Base To Base				G6
_		165-8	QC 750	Quality Control		⊗		1.00		3,4		5.57	Base To Base			L	D <b>7</b>
2	16-534006g\16-0783.wiff		QC 10000	Quality Control		☒			9624.3	-3.8		5.59	Base To Base				47
1	16-534006g\16-0764.wiff		Mobile Phase	Unknown						WA			No Peak				D1
_			46682, 6-12 hr	Unknown						#BAD!		5.52	Base To Base		***************************************		D1
			46682, 6-12 hr rinse 46683, 6-12 hr	Unknown Unknown			N/A N/A		671,40 No intercept	N/A #BAD!		5.50 5.53	Base To Base	<u> </u>		; ·	D1 .
	16-534006g\16-0788.wilf 16-534006g\16-0788.wilf			Unknown					No intercept 3042.0	#BAD!		5.53 5.44	Base To Base	<u> </u>			D1
			46690, 6-12 hr	Unknown				1.00	No Intercept	#BAD!		5.44 5.55	Base To Base Base To Base				D1
2			46690, 6-12 hr rinse	Unknown					1813,9	N/A		5.51	Base To Base	<del>                                     </del>			D1
		168-1	46664, 12-24 hr	Unknown			N/A		No Intercept	#BAD!		5.53	Base To Base				D1
-	16-534006g\16-0792.wilf		46664, 12-24 hr rinse	Unknown				1,00	752.56	N/A		5.46	Base To Base			i	D1
Š	16-534006g\16-0793,wiff		46667, 12-24 hr	Unknown						#BAD!		5.57	Base To Base	┝┼			D1
Ī	16-534006g\16-0794.wilf		46667, 12-24 hr rinse	Unknown					1390.8	N/Λ		5.46	Base To Base				D1
		168-5	46670, 12-24 hr	Unknown				1.00	No Intercept	#BAD!		5.59	Base To Base				Ď1
ँ	16-534006g\16-0796.wiff		46670, 12-24 hr rinse	Unknown	_				632.48	N/A		5.47	Valley	ă		10	D1
	16-534006q\16-0797,wiff		46682, 12-24 hr	Unknown			N/A	1.00	1258.1	N/A	1.76e+007	5.61	Base To Base			10	

Printing Date: Thursday, March 22, 2007

Operator: Shelley Hollar

	File Name	Sample ID	Sample Name	Sample Type	Analyte Peal Name	Use Record	Analyte Concentration (ng/mL)	Dilution Factor	Calculated Concentration (ng/mL)	%RE	Analyte Peak Area (counts)	Analyte Retention Time (min)	Analyte Integration Type	Record Modified	Sample Annotation	LCARTRIDGET RAY	LCARTRIDGEP OSITION
66	16-534006g\16-0798.wiff	168-8	46682, 12-24 hr rinse	Unknown			N/A	1.00	238.01	N/A	3.65e+006	5.51	Base To Base	П	•	10	ID1
67	16-534006g\16-0799.wiff	168-9	46683, 12-24 hr	Unknown		1	N/A	1.00	1100.8	N/A	1.56e+007	5.58	Base To Base			10	D1
	16-534006g\16-0800,wilf		46683, 12-24 hr rinse	Unknown	_	1	N/A	1.00	342.42	N/A	5.19e+006	5.48	Base To Base			10	D1
69	16-534006g\16-0801.wilf	168-11	46690, 12-24 hr	Unknown		1	N/A	1.00	616.33	N/A	9.10e+006	5.60	Base To Base			10	D1
70	16-534006g\16-0802.wiff	168-12	46690, 12-24 hr rinse	Unknown		1	N/A	1.00	122,72	N/A	1.94e+006	5.49	Base To Base			10	D1
	16-534006g\16-0803.wiff		Mobile Phase	Unknown		1	N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			10	C6
72	16-534006g\16-0804.wiff	165-3	QC 30	Quality Control	-	⊠	30.000	1.00	28.295	-5.7	5.11e+005	5.58	Base To Base	h	·····	10	E6
	16-534006g\16-0805.wiff		QC 250	Quality Control		⊠	250.00	1.00	217.17	-13.	3.35e+006	5.65	Base To Base		***************************************	10	G6
	16-534006g\16-0806.wiff		QC 750	Quality Control		⊠	750.00	1.00	755.20	0.69	1.10e+007	5.65	Base To Base			10	D7
75	16-534006g\16-0807.wiff	165-12	QC 10000	Quality Control	_	(⊠)	10000.	20,0	9656.7	-3.4	7.21e+006	5.56	Base To Base			10	A7
76	16-534006g\16-0808,wiff	168a-1	Mobile Phase	Unknown		Ì	N/A	1.00	No Peak	NΑ	0.00e+000	0.00	No Peak		······································	10	D1
77	16-534006g\16-0809.wiff	164-3	C 10	Standard		<b> </b>	10.000	1.00	8.4934	-15.	2.08e+005	5.46	Base To Base		***************************************	10	E3
78	16-534006g\16-0810,wiff	164-6	C 30	Standard		Ø	30.000	1.00	27.228	-9.2	4.94e+005	5.62	Base To Base	- H	······································		F3
79	IG-534006g\IG-0811.wiff	164-9	C 100	Standard	_	8	100.00	1.00	100.59	0.59	1.61e+006	5.61	Base To Base		······	10	B4
80	16-534006g\16-0812.will	164-12	C 300	Standard		×	300.00	1.00	293.97	-2.0	4.48e+006	5.68	Base To Base			10	E4
81	16-534006g\16-0813.will	164-15	C 1000	Standard		×	1000.0	1,00	960.64	-3.9	1.38e+007	5.56	Base To Base				G4





Printing Date: Thursday, March 22, 2007

Operator: Shelley Hollar

-122

218 of 281

Note:

Experimental serum samples

Study Record Page: 201a

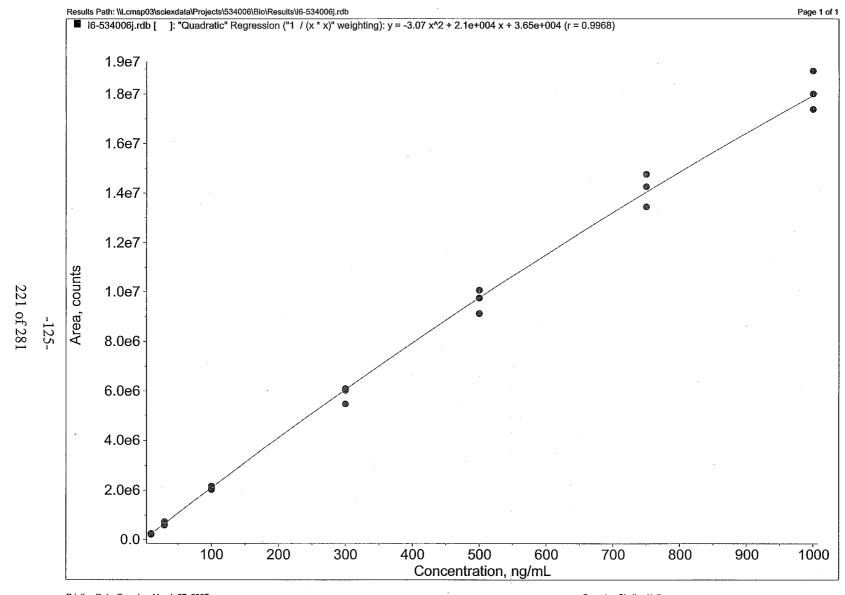
219	
of 281	-123-

Resu	Its Path: \\Lcmsp03	sciexda	ta\Projects\534006\	Bio\Results\I6	5-534006j.rd	b										Page 1 of
	File Name	Sample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record	Analyte Concentration (ng/mL)	Dilution Factor	Calculated Concentration (ng/mL)	%RE	Analyte Peak Area (counts)	Analyte Retention Time (min)	Analyte Integration Type	Record Modified	Sample Annotation	
	l6-534006j\l6-1018.wiff		Test Solution	Unknown			N/A	1.00	1470.8	N/A	2.42e+007	5.81	Base To Base			
	16-534006j\16-1019.wiff		Test Solution	Unknown			N/A	1.00	1512.7	N/A	2.47e+007	5.70	Base To Base	⊠	peak tailing factor	
3	16-534006j\16-1020.wiff		Test Solution	Unknown			N/A	1.00	1583,3	N/A	2.55e+007	5.69	Valley			
1	16-534006j\16-1021.wiff		Test Solution	Unknown			N/A	1.00	1660.4	N/A	2.64e+007	5.62	Base To Base			1
5	16-534006j\16-1022.wiff		Test Solution	Unknown			N/A	1.00	1739.0	N/A	2.72e+007	5.63	Base To Base			
5	16-534006j\l6-1023.wiff 16-534006j\l6-1024.wiff		Test Solution Test Solution	Unknown			N/A N/A	1.00	1872.2	N/A	2.85e+007	5.57	Base To Base	<u> </u>		
ļ	16-534006j\16-1025.wiff		Test Solution					1.00	1882.1	N/A	2.86e+007	5.54	Base To Base	<u> </u>		
	16-534006j\16-1025.wilf		Test Solution	Unknown Unknown			N/A N/A	1,00	1969.0 2088.8	N/A N/A	2.94e+007	5.51	Base To Base			
	16-534006j\16-1027.will	J	Test Solution	Unknown			N/A	1.00	2186.0	N/A	3.04e+007 3.12e+007	5.45 5.42	Base To Base		peak tailing factor	- Ingeres
	16-534006j\16-1028.wiff	1	Mobile Phase	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	Base To Base No Peak	<u> </u>		
	16-534006j\16-1029.wiff	I	Solvent Blank	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			
13	16-534006j\16-1030.wiff	L	Blank Serum	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			
	16-534006j\l6-1031.wiff		Blank Serum	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak	<u> </u>		
100 1 1 100	16-534006j\16-1032.will	1	Blank Serum	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			
16	16-534006j\16-1033.wilf		Mobile Phase	Unknown			N/A	1.00	No Peak	NA	0.00e+000	0.00	No Peak		······	
	16-534006j\l6-1034.wiff		C 10	Standard	*******		10.000	1.00	8.6731	-13,	2.18e+005	5.13	Base To Base	- -		
	l6-534006j\l6-1035.wiff		C 30	Standard	******		30.000	1.00	26.882	-10.	5.98e+005	4.94	Base To Base			
19	16-534006j\16-1036.wiff	192-7	C 100	Standard	*******		100.00	1.00	96.152	-3.8	2.03e+006	4.97	Base To Base			
20	16-534006j\16-1037.wiff	192-10	C 300	Standard	*******		300.00	1.00	269.80	-10.	5.47e+006	5.01	Base To Base			
21	16-534006j\16-1038.wiff	192-13	C 500	Standard			500.00	1.00	464.53	-7.1	9.12e+006	4.95	Base To Base			·
22	l6-534006j\l6-1039.wiff	192-16	C 750	Standard			750.00	1.00	713.59	-4.9	1.34e+007	4.89	Base To Base			
	16-534006j\16-1040.wiff	192-19	C 1000	Standard			1000.0	1.00	963.22	-3.7	1.74e+007	4.87	Base To Base			
	16-534006j\16-1041.will	188-1	Mobile Phase	Unknown	********		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			
25	16-534006j\16-1042.wilf	193-1	QC 30	Quality Control		⊠	30,000	1.00	28.320	-5.6	6.28e+005	4.91	Valley		***************************************	
	16-534006j\16-1043.wiff		QC 250	Quality Control			250.00	1.00	235.59	-5.8	4.81e+006	4.90	Base To Base			
	16-534006j\16-1044.wiff		QC 750	Quality Control			750.00	1.00	732.32	-2.4	1.38e+007	4.93	Base To Base			
	16-534006j\16-1045.wiff		QC 30000	Quality Control			30000.	1000.	27980.	-6.7	6.21e+005	4.84	Base To Base			
	16-534006j\16-1046.wiff		Mobile Phase	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			
	l6-534006j\l6-1047.wiff		46662,D0,1M,T0	Unknown	<u></u>		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			
	16-534006j\16-1048.wiff		46663,D0,1M,T0	Unknown			N/A	1.00	No Peak	N/A	0.00e+000		No Peak			
	16-534006j\16-1049.wiff		46665,D0,1M,T0	Unknown			N/A		No Peak	N/A	0.00e+000	0.00	No Peak			
	16-534006j\16-1050,wiff		46678,D0,1F,T0	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		······	
7.000000	l6-534006j\l6-1051.wilf l6-534006j\l6-1052.wilf		46679,D0,1F,T0	Unknown			N/A	1.00	No Peak	N/A	0.00e+000		No Peak			
	16-534006j\l6-1053,wiff		46680,D0,1F,T0 46666,D0,1M,T2 min	Unknown			N/A N/A	1.00 1000.	No Peak 85509.	N/A	0.00e+000		No Peak			
A 7 A	16-534006j\16-1054.wiff			Unknown			N/A	1000.	16736.	N/A N/A	1.81e+006 3.87e+005	ž	Base To Base			
	16-534006j\l6-1055.wiff		46672,D0,1M,T2 min	Unknown			N/A	1000.	100720.		2.12e+006	4.78 4.84	Base To Base			
	16-534006j\16-1056.wiff		46681,D0,1F,T2 min	Unknown			N/A	I	88738.	N/A	1.87e+006	Ł	Base To Base Base To Base			
	I6-534006j\I6-1057.wiff		46684,D0,1F,T2 min	Unknown					98125.	N/A	2.07e+006		Base To Base			
3.00	16-534006j\16-1058.wiff		46685.D0,1F,T2 min	Unknown			N/A		97362.	N/A	2.05e+006		Valley		·	
	16-534006j\l6-1059.wiff	l	46673,D0,1M,T10min	Unknown			N/A		71736.	N/A	1.53e+006		Base To Base			
	16-534006j\16-1060.wiff		46674,D0,1M,T10min	Unknown			N/A		59438.	N/A	1.27e+006	4.78	Base To Base			
	16-534006j\16-1061.wiff		46676,D0,1M,T10min	Unknown			N/A		76212.	N/A	1.62e+006		Base To Base	┝╫┪		
45	16-534006j\16-1062.wiff	196-10	46686,D0,1F,T10min	Unknown			N/A	1000.	53064.	N/A	1.14e+006	\$	Base To Base			
	16-534006j\16-1063.wiff		46688,D0,1F,T10min	Unknown	-		N/A	1000.	43359.	N/A	9.40e+005		Base To Base			
47	16-534006j\16-1064.wiff		46691,D0,1F,T10min	Unknown					54242.	N/A	1.17e+006		Base To Base			
48	l6-534006j\l6-1065.wiff		46662,D0,1M,T20 min	Unknown			N/A		65371.		1.39e+006		Base To Base	버		
49	16-534006j\16-1066.wilf	196-14	46663,D0,1M,T20 min	Unknown			N/A	1000.	55817.	N/A	1.20e+006	4.75	Base To Base	i i		
	16-534006j\16-1067.wiff	196-15	46665,D0,1M,T20 min	Unknown			N/A	1000.	65870.	N/A	1.41e+006		Base To Base			
	16-534006j\l6-1068.wilf	196-16	46678,D0,1F,T20 min	Unknown			N/A	1000.	30804.	N/A	6.80e+005		Base To Base		peak lailing factor	
	l6-534006j\l6-1069.wiff		46679,D0,1F,T20 min	Unknown			N/A	1000.	27712.	N/A	6.16e+005	4.72	Base To Base			
	16-534006j\16-1070.wiff		46680,D0,1F,T20 min	Unknown	•			1000.	31506.	N/A	6.94e+005	4.74	Base To Base			
	l6-534006j\l6-1071.wiff		Mobile Phase	Unknown			N/A		No Peak	N/A	0.00e+000	0.00	No Peak			
	16-534006j\16-1072.wiff		C 10	Standard				1.00	10.445	4.5	2.55e+005	4.75	Base To Base		***************************************	
	l6-534006j\l6-1073.wiff		C 30	Standard		×		1.00	33.657	12.	7.39e+005		Base To Base			
	l6-534006j\l6-1074.wiff		C 100	Standard			100.00		99.328	-0.67	2.09e+006		Base To Base			
58	l6-534006j\l6-1075.wiff	192-11	C 300	Slandard		☒	300.00	1.00	298.41	-0.53	6.02e+006	4.66	Base To Base			

Printing Date: Tuesday, March 27, 2007

Operator: Shelley Hollar

	File Name	Sample ID	Sample Name	Sample Type	Analyle Peak Name	Use Record	Analyte Concentration (ng/mL)	Dilution Factor	Calculated Concentration (ng/mL)	%RE	Analyte Peak Area (counts)	Analyte Retention Time (min)	Analyte Integration Type	Record Modified	Sample Annotation
X	16-534006j\16-1076.wiff	192-14	C 500	Standard		⊠	500.00	1.00	499.45	-0.11	9.75e+006	4.69	Base To Base		**************************************
jk	16-534006j\l6-1077.wiff	192-17	C 750	Standard	,,,,,	⊠	750.00	1.00	763.47	1.8	1.43e+007	4.63	Base To Base		
	16-534006j\l6-1078.wiff	192-20	C 1000	Standard		⊠	1000.0	1.00	1004.9	0.49	1.80e+007	4.67	Base To Base		***************************************
V	16-534006j\16-1079.will	188-1	Mobile Phase	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
8.	16-534006j\16-1080.wiff	193-2	QC 30	Quality Control	· ·	⊠	30.000	1.00	30.544	1.8	6.74e+005	4.67	Base To Base		
	16-534006j\16-1081.wiff	193-5	QC 250	Quality Control		☒	250.00	1.00	242.87	-2.9	4.95e+006	4.69	Base To Base	П	
	16-534006j\16-1082.wiff	193-8	QC 750	Quality Control		⊠	750.00	1.00	735.43	-1.9	1.38e+007	4.65	Base To Base		
33	16-534006j\16-1083.wiff	193-11	QC 30000	Quality Control		⊠	30000,	1000.	30449.	1.5	6.72e+005	4.66	Base To Base		***************************************
	16-534006j\16-1084.wiff	188-1	Mobile Phase	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
	16-534006j\16-1085.wiff	197-1	46666,D0,1M,T30 min	Unknown			N/A	1000.	51702.	N/A	1.11e+006	4.64	Base To Base		
ď	16-534006j\16-1086.wiff	197-2	46669,D0,1M,T30 min	Unknown			N/A	1000.	19975.	N/A	4.54e+005	4.68	Base To Base		***************************************
V.	16-534006j\16-1087.wiff	197-3	46672,D0,1M,T30 min	Unknown			N/A	1000.	57560.	N/A	1.23e+006	4.73	Base To Base		
V.	16-534006j\16-1088.will	197-4	46681,D0,1F,T30 min	Unknown			N/A	1000.	27329.	N/A	6.08e+005	4.66	Base To Base		······································
¢	16-534006j\16-1089.wiff	197-5	46684,D0,1F,T30 min	Unknown			N/A	1000.	24091.	N/A	5.40e+005	4.76	Base To Base		
	16-534006j\16-1090.wiff	197-6	46685,D0,1F,T30 min	Unknown			N/A	1000.	19724.	N/A	4.49e+005	4.68	Base To Base		
3	16-534006j\l6-1091,wiff	197-7	46673,D0,1M,T1 hr	Unknown			N/A	1000.	62374.	N/A	1.33e+006	4.72	Base To Base		····
	16-534006j\16-1092.wiff	197-8	46674,D0,1M,T1 hr	Unknown			N/A	1000.	47426.	N/A	1.02e+006	4.70	Base To Base		***************************************
	16-534006j\l6-1093.will	197-9	46676,D0,1M,T1 hr	Unknown			N/A	1000.	81376.	N/A	1.72e+006		Base To Base		
	16-534006j\l6-1094.wiff	197-10	46686,D0,1F,T1 hr	Unknown			N/A	1000.	15651.	N/A	3.64e+005		Base To Base		
S	16-534006j\16-1095.wiff	197-11	46688,D0,1F,T1 hr	Unknown			N/A	1000.	7912.2	WA	2.02e+005	4.67	Base To Base		**************************************
	16-534006j\16-1096.wiff	197-12	46691,D0,1F,T1 hr	Unknown			N/A	1000.	17595.	N/A	4.05e+005	!	Base To Base		
1	16-534006j\l6-1097.wiff	198-1	46662,D0,1M,T3 hr	Unknown			N/A	100.	37375.	N/A	7.45e+006		Base To Base		······································
8	16-534006j\16-1098.wiff	198-2	46663,D0,1M,T3 hr	Unknown			N/A	100.	15238.	N/A	3.16e+006		Base To Base		
	16-534006j\16-1099.wiff	198-3	46665,D0,1M,T3 hr	Unknown		***************	N/A	100.	22245.	N/A	4.55e+006		Base To Base		
	i6-534006j\l6-1100.wiff	198-4	46678,D0,1F,T3 hr	Unknown			N/A	100.	1028.0	N/A	2.52e+005	4.62	Base To Base		
	16-534006j\16-1101.wiff	198-5	46666,D0,1M,T5 hr	Unknown			N/A	100.	8626.8	N/A	1.82e+006	4.61	Base To Base	H	······································
7	16-534006j\16-1102.wiff	198-6	46669,D0,1M,T5 hr	Unknown			N/A	100.	20677.	N/A	4.24e+006		Base To Base		
	16-534006j\l6-1103.wiff	198-7	46672,D0,1M,T5 hr	Unknown	-		N/A	100.	17947.	N/A	3.70e+006	4.69	Base To Base	T T	
	16-534006j\16-1104.wiff	198-8	46673,D0,1M,T7 hr	Unknown			N/A	100.	24227.	N/A	4.94e+006		Base To Base		
7.	16-534006j\16-1105.wiff	198-9	46674,D0,1M,T7 hr	Unknown			N/A	100.	7111.1	N/A	1.51e+006	4.61	Base To Base	H	******
	16-534006j\16-1106.wiff	198-10	46676,D0,1M,T7 hr	Unknown			N/A	100.	20778.	N/A	4.26e+006	4.64	Base To Base		
	16-534006j\16-1107.wiff	201-2	Stock Stab	Unknown			N/A	1.00	1662.6	N/A	2.64e+007	4.68	Base To Base	H	
F	16-534006j\16-1108.wiff	201-4	Stock Stab	Unknown			N/A	1.00	1636.1	N/A			Base To Base		
"	16-534006j\16-1109.wiff	201-6	Stock Stab	Unknown			N/A	1.00	1661.7	N/A	2.64e+007		Base To Base		······································
	16-534006j\16-1110.wilf	201-8	Slock Stab	Unknown			N/A	1.00	1683.4	N/A	2.66e+007		Base To Base	H	······································
	l6-534006j\l6-1111.wiff	188-1	Mobile Phase	Unknown			N/A	1.00	No Peak	N/A			No Peak	H	
	16-534006j\16-1112.wiff	193-3	QC 30	Quality Control		⊠	30,000	1.00	30.124	0.41			Base To Base	H	
	16-534006j\16-1113.wiff	193-6	QC 250	Quality Control		\(\overline{\pi}\)	250.00	1.00	248.88	-0.45		L	Base To Base		
	16-534006j\16-1114.wiff	193-9	QC 750	Quality Control		Ø	750.00		742.94	-0.94			Base To Base	H	· · · · · · · · · · · · · · · · · · ·
	l6-534006j\l6-1115.wiff	193-12	QC 30000	Quality Control		×	30000.	1000.	30268.	0.89	1		Base To Base	H	
Š.	16-534006j\16-1116.wiff	188-1	Mobile Phase	Unknown			N/A	1.00	No Peak	N/A	l		No Peak		
	16-534006j\16-1117.wiff	192-3	C 10	Standard		⊠	10.000	1.00	10.500	5.0			Base To Base		
	i6-534006j\l6-1118.wiff	192-6	C 30	Standard		- ×	30.000	1.00	33.321	11.			Base To Base		
	l6-534006j\l6-1119.wiff	192-9	C 100	Standard		☒	100.00	1.00	103.22	3.2			Base To Base		
	16-534006j\16-1120.wiff	192-12	C 300	Standard		<u> </u>	300.00	1.00	302.10	0.70	ł		Base To Base	H	
-	16-534006j\16-1121.wiff	192-15	C 500	Standard		×	500.00		517,14	3.4	<u> </u>		Base To Base		~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~
Ť	16-534006j\16-1122,wiff	192-18	C 750	Standard		⊠	750.00	1.00	794.40	5.9	L		Base To Base	H	
3		1	C 1000	Standard		☒	1000,0	1.00	1068.8	6.9			Base To Base	-H	······································



Printing Date: Tuesday, March 27, 2007

Operator: Shelley Hollar

222 of 281 -126-

Note:

Experimental serum sample analysis

11-day long term frozen stability and 4 hour benchtop stability assessments

Study Record Page: 212a

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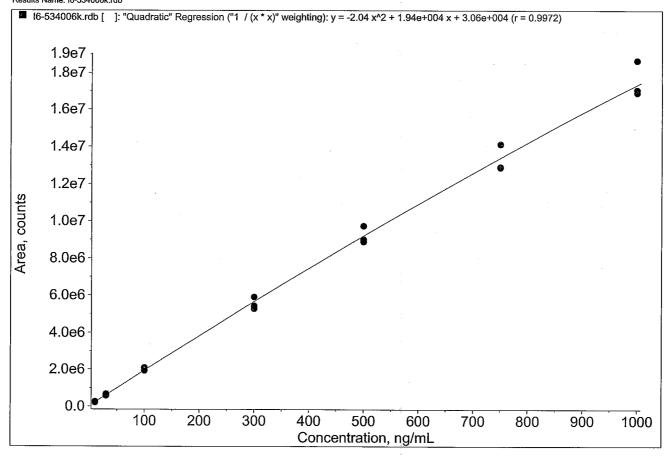
223 of 281

			Fig. 2 Jan St. Jan St. St. St. St. St. St. St. St. St. St.		C00002305.350000			1		1	100000000000000000000000000000000000000	I	I	Tessessia		
	File Name	Sample ID	Sample Name	Sample Type	Analyle Peak Name	Use Record	Analyte Concentration (ng/mL)	Dilution Factor	Calculated Concentration (ng/mL)	%RE	Analyte Peak Area (counts)	Analyte Retention Time (min)	Analyte Integration Type	Record Modified	Sample Annotation	
	16-534006k\16-1124.will	203-10	Test Solution	Unknown		LAST PERSON	N/A	1.00	2391.2	N/A	3,47e+007	5.27	Base To Base			
×.	l6-534006k\l6-1125.wiff	203-10	Test Solution	Unknown			N/A	1.00	2494.8	N/A	3.56e+007	5.37	Base To Base		***************************************	
	16-534006k\16-1126,wiff	203-10	Test Solution	Unknown	ļ		N/A	1.00	2510.0	N/A	3.58e+007	5.41	Base To Base			
100		203-10	Test Solution	Unknown			N/A	1.00	2594.9	N/A	3.65e+007	5.43	Base To Base			
άij,		203-10	Test Solution	Unknown	Ī		N/A	1.00	2631.4	N/A	3.69e+007	5.44	Base To Base		***************************************	
V.		203-10	Test Solution	Unknown	Ī		N/A	1.00	2645.6	N/A	3.70e+007	5.45	Base To Base			
	l6-534006k\l6-1130.wiff	203-10	Test Solution	Unknown	i		N/A	1.00	2687.3	N/A	3.73e+007	5.49	Base To Base		······································	
	l6-534006k\l6-1131.wiff	203-10	Test Solution	Unknown	·		N/A	1.00	2778.9	N/A	3.81e+007	5.60	Base To Base			
269	I6-534006k\I6-1132.wiff	203-10	Test Solution	Unknown			N/A	1.00	2817.1	N/A	3.84e+007	5.52	Base To Base			
) .	l6-534006k\l6-1133.wiff	203-10	Test Solution	Unknown	Ī		N/A	1.00	2793.5	N/A	3.82e+007	5.55	Base To Base			
1	l6-534006k\l6-1134.wiff		Mobile Phase	Unknown	Î		N/A	1.00	No Peak	N∕A	0.00e+000	0.00	No Peak		······································	
2		207-1	Solvent Blank	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		**************************************	
3	I6-534006k\I6-1136.wiff	207-2	Blank Serum	Unknown	T		N/A	1.00	0.58423	N/A	4.19e+004 ·	5.34	Base To Base		······································	
1		207-3	Blank Serum	Unknown			N/A	1.00	0.43492	N/A	3.90e+004	5.49	Base To Base			
5		207-4	Blank Serum	Unknown			N/A	1.00	0.48626	N/A	4.00e+004	5.33	Base To Base			
3	16-534006k\16-1139.wiff		Mobile Phase	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			
7			C 10	Standard		☒	10.000	1.00	9.6359	-3.6	2.17e+005	5.51	Base To Base		······································	
8	l6-534006k\l6-1141.wiff	205-4	C 30	Standard	Ī	⊠	30.000	1.00	29.916	-0.28	6.08e+005	5.42	Base To Base			
9	l6-534006k\l6-1142.wiff	205-7	C 100	Standard		⊠	100.00	1.00	98.972	-1.0	1.93e+006	5.41	Base To Base		······································	
0	l6-534006k\l6-1143.wilf	205-10	C 300	Standard		⊠	300,00	1.00	286.86	-4.4	5.41e+006	5.39	Valley		······································	,
1	l6-534006k\l6-1144.wiff	205-13	C 500	Standard		⊠	500.00	1.00	488.28	-2.3	8.99e+006	5.40	Base To Base		***************************************	
2			C 750	Standard		⊠	750.00	1.00	719.77	-4.0	1.29e+007	5.38	Base To Base			
3	l6-534006k\l6-1146.wiff	205-19	C 1000	Standard	ſ	⊠	1000.0	1.00	981.41	-1.9	1.71e+007	5.42	Base To Base		***************************************	
4 :	l6-534006k\l6-1147.wiff		Mobile Phase	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		***************************************	
5	l6-534006k\l6-1148.wiff	206-1	QC 30	Quality Control		⊠	30,000	1.00	30.757	2.5	6.24e+005	5.44	Base To Base	T n i		
5	l6-534006k\l6-1149.wiff	206-4	QC 250	Quality Control		⊠	250.00	1.00	245.68	-1.7	4.66e+006	5.37	Base To Base		······································	
7	l6-534006k\l6-1150.wiff	206-7	QC 750	Quality Control	******	⊠	750.00	1.00	734.37	-2,1	1.31e+007	5.39	Base To Base			
8 .	l6-534006k\l6-1151.wiff	206-10	QC 30000	Quality Control		⊠	30000.	100.	29723.	-0.92	5.60e+006	5.36	Base To Base			
9	l6-534006k\l6-1152.wilf		Mobile Phase	Unknown			N/A	1.00	< 0	#BAD!	1.43e+004	5.43	Base To Base		······································	
0	l6-534006k\l6-1153.wiff	208-1	46688,D0,1F,T1	Unknown			N/A	100.	10027.	N/A	1.95e+006	5.35	Base To Base			
1	16-534006k\16-1154.wiff		Mobile Phase	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak	T 🗖 T		
2		209-1	LT Stb 30 ng/mL	Unknown			N/A	1.00	30.493	N/A	6.19e+005	5.38	Base To Base		· · · · · · · · · · · · · · · · · · ·	
3	16-534006k\16-1156.wiff	209-2	LT Slb 30 ng/mL	Unknown			N/A	1.00	31.482	N/A	6.38e+005	5.36	Base To Base			
4	16-534006k\16-1157.wilf	209-3	LT Stb 30 ng/mL	Unknown			N/A	1.00	30.472	N/A	6.18e+005	5.33	Base To Base			
5		209-4	LT Stb 750 ng/mL	Unknown			N/A	1.00	772.64	N/A	1.38e+007	5.33	Base To Base			
6	16-534006k\16-1159.wiff	209-5	LT Stb 750 ng/mL	Unknown			N/A	1.00	759.55	N/A	1.36e+007	5.34	Base To Base			
7.	l6-534006k\l6-1160.wiff	209-6	LT Stb 750 ng/mL	Unknown			N/A	1.00	763.98	N/A	1.36e+007	5.39	Base To Base			
8	16-534006k\16-1161.wiff		Mobile Phase	Unknown	-		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			
9	l6-534006k\l6-1162.wiff	205-2	C 10	Standard		×	10.000	1.00	8.9138	-11.	2.03e+005	5.33	Base To Base			
0	16-534006k\16-1163.wiff	205-5	C 30	Standard		×	30.000	1.00	27.780	-7.4	5.67e+005	5.30	Base To Base	l fil	***************************************	
1	16-534006k\16-1164.wiff	205-8	C 100	Standard		×	100.00	1.00	98.612	-1.4	1.92e+006	5.34	Base To Base			
2	l6-534006k\l6-1165.wiff	205-11	C 300	Standard		⊠	300.00	1.00	278.75	-7.1	5.27e+006	5.28	Base To Base			
3	16-534006k\16-1166,wiff	205-14	C 500	Slandard	******	×	500.00	1.00	481.28	-3.7	8.87e+006	5,33	Base To Base			
4	l6-534006k\l6-1167.wiff	205-17	C 750	Slandard			750.00	1.00	718.19	-4.2	1.29e+007	5.27	Base To Base		~~~~~~	
5	l6-534006k\l6-1168.wilf	205-20	C 1000	Standard		×	1000.0	1.00	970.81	-2.9	1.69e+007	5.31	Base To Base			
8	16-534006k\16-1169.wiff	-	Mobile Phase	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			
7	I6-534006k\l6-1170.wiff	206-2	QC 30	Quality Control		×	30.000	1.00	30,556	1.9	6.20e+005	5.26	Base To Base			
3	16-534006k\16-1171.wiff	206-5	QC 250	Quality Control		× ×	250.00	1.00	242.13	-3.1	4.60e+006	5.30	Base To Base			
)	I6-534006k\I6-1172.wiff	206-8	QC 750	Quality Control	• • • • • • • • • • • • • • • • • • • •		750.00	1,00	746.88	-0.42	1.33e+007	5.33	Base To Base			
)	l6-534006k\l6-1173.wilf		QC 30000	Quality Control			30000.	100.	28014.	-6.6	5.29e+006	5.29	Base To Base			
1	16-534006k\16-1174.wiff		Mobile Phase	Unknown		<del></del>	N/A	1.00	< 0	#BAD!	1.45e+004	5.25	Base To Base	$\vdash \vdash \vdash$	•••••••••••••••••••••••••••••••••••••••	
2	l6-534006k\l6-1175.wiff	210-1	4 Hr Slb 30 ng/mL	Unknown			N/A	1.00	30.236	N/A	6.14e+005		Base To Base	f + f		
			4 Hr Stb 30 ng/mL	Unknown			N/A		30.396	N/A	6.17e+005		Base To Base			
			4 Hr Stb 30 ng/mL	Unknown	· —		N/A	1.00	30.575	N/A	6.20e+005		Base To Base	ᅡ┼		
5			4 Hr Stb 750 ng/mL	Unknown			N/A	1.00	762.58	N/A	1.36e+007		Base To Base			
5	l6-534006k\l6-1179.wilf		4 Hr Stb 750 ng/mL	Unknown			N/A	1.00	772.25	N/A	1,38e+007		Base To Base	$\vdash \exists \vdash$		
21.7.1			4 Hr Slb 750 ng/ml.	Unknown			N/A		755.26	N/A	1.35e+007		Base To Base	$\vdash$		
3.3.	l6-534006k\l6-1181.wiff		Mobile Phase	Unknown			N/A	1.00	No Peak		1		No Peak			

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	File Name	Sample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record	Analyte Concentration (ng/mL)	Dilution Factor	Calculated Concentration (ng/mL)	%RE	Analyte Peak Area (counts)	Analyte Retention Time (mln)	Analyte Integration Type	Record Modified	Sample Annotation
	l6-534006k\l6-1182.wiff		QC 30	Unknown	1000000	1	N/A	1.00	38.134	N/A	7.66e+005	5.36	Base To Base		
0	16-534006k\16-1183.wiff		QC 30	Unknown		1	N/A	1.00	36.120	N/A	7.27e+005	5.36	Base To Base		
1	16-534006k\16-1184.wiff		QC 30	Unknown			N/A	1.00	35.753	N/A	7.20e+005	5.42	Base To Base		
2	16-534006k\16-1185.wiff		QC 250	Unknown		Î	N/A·	1.00	301.70	N/A	5.68e+006	5.38	Base To Base		
	I6-534006k\I6-1186.wiff		QC 250	Unknown		Ī	N/A	1.00	297.31	N/A	5.60e+006	5.35	Base To Base		
	l6-534006k\l6-1187.wiff		QC 250	Unknown	1		N/A	1.00	293.31	N/A	5.53e+006	5.30	Base To Base		***************************************
	16-534006k\16-1188.wiff		QC 750	Unknown			N/A	1.00	884.01	N/A	1.55e+007	5.35	Base To Base		
6	16-534006k\16-1189.wilf		QC 750	Unknown	l		N/A	1.00	877.09	N/A	1.54e+007	5.29	Base To Base		***************************************
7 🕢	16-534006k\16-1190.wiff		QC 750	Unknown			N/A	1.00	868.73	N/A	1.53e+007	5.23	Base To Base		
В	l6-534006k\l6-1191.wiff	193-10	QC 30000	Unknown			N/A	1000.	35970.	N/A	7.24e+005	5.27	Base To Base		
9 💥	l6-534006k\l6-1192.wiff	193-11	QC 30000	Unknown		Ī	N/A	1000.	37506.	N/A	7.54e+005	5.28	Base To Base	Ħ	
0	6-534006k\ 6-1193.wiff	193-12	QC 30000	Unknown	,,,,,,		N/A	1000.	37151.	N∕A	7.47e+005	5,30	Base To Base	T I	
1	16-534006k\16-1194,wiff		Mobile Phase	Unknown			N/A	1.00	< 0	#BAD!	1.71e+004	5.35	Base To Base	ᆔᅥ	
2	16-534006k\16-1195,wiff	155-1	QC 30	Unknown	,		N/A	1.00	166.76	N/A	3.20e+006	5.31	Base To Base		***************************************
3	16-534006k\16-1196.wiff	155-2	QC 30	Unknown			N/A	1.00	67.131	N/A	1.32e+006		Base To Base		
	l6-534006k\l6-1197,wiff	155-3	QC 30	Unknown			N/A	1.00	62.723	N/A			Base To Base		<del></del>
5	16-534006k\l6-1198.will	155-4	QC 250	Unknown	*****	l	N/A	1.00	535.08	N/A			Base To Base		····
3	16-534006k\16-1199.wiff	155-5	QC 250	Unknown		l	N/A	1.00	492.76	N/A	9.07e+006		Base To Base		***************************************
7.55	16-534006k\l6-1200.wiff	155-6	QC 250	Unknown			N/A	1.00	540.93	N/A		5.37	Base To Base		······
3	l6-534006k\l6-1201,wiff	155-7	QC 750	Unknown			N/A	1.00	1721.7	N/A	2.73e+007	5.32	Base To Base		
)	16-534006k\l6-1202.wiff	155-8	QC 750	Unknown			N/A	1.00	1794.3	N/A	2.82e+007		Base To Base		
)	l6-534006k\l6-1203.wilf	155-9	QC 750	Unknown			N/A	1.00	1707.3	N/A	2.71e+007		Base To Base	౼౼	
i je	16-534006k\16-1204.vviff	155-10	QC 10000	Unknown	,		N/A	20.0	21073.	N/A			Base To Base		
2	16-534006k\16-1205.wiff	155-11	QC 10000	Unknown	******		N/A	20.0	23424.	N/A	1.99e+007		Base To Base		
3	16-534006k\16-1206.wiff	155-12	QC 10000	Unknown			N/A	20.0	23338.	N/A			Base To Base		
	l6-534006k\l6-1207.wiff		Mobile Phase	Unknown	******		N/A	1.00	No Peak	N/A	0.00e+000		No Peak	ㅠĦㅠ	
5	l6-534006k\l6-1208.wilf	206-3	QC 30	Quality Control	*******	⊠	30.000	1.00	32.445	8.2			Base To Base	-51	
5	16-534006k\16-1209.wiff	206-6	QC 250	Quality Control		Ø	250.00	1.00	260.75	4.3			Base To Base		·····
	16-534006k\16-1210.wiff	206-9	QC 750	Quality Control			750.00	1.00	791.55	5.5	·		Base To Base		
5.0	l6-534006k\l6-1211.wiff	206-12	QC 30000	Quality Control		冈	30000.	100.	33381.	11.	L		Base To Base	-6-	
1	l6-534006k\l6-1212.wiff		Mobile Phase	Unknown	********		N/A	1.00		N/A	L		No Peak		
)	l6-534006k\l6-1213,wiff	205-3	C 10	Standard		Ø	10.000	1.00		14.			Base To Base		
	l6-534006k\l6-1214.wilf	205-6	C 30	Standard	******		30.000			·			Base To Base	片	
2	l6-534006k\l6-1215.wiff	205-9	C 100	Standard						7.2			Base To Base	-H	***************************************
1	16-534006k\16-1216.wiff	205-12	C 300	Standard			300.00	1.00		4.4			Base To Base		
	16-534006k\16-1217.wiff	205-15	C 500	Slandard			500.00						Base To Base		
	16-534006k\16-1218.wiff	205-18	C 750	Standard			750.00						Base To Base	-H	
	16-534006k\16-1219.wiff	205-21	C 1000	Standard									Base To Base		



Printing Date: Wednesday, March 28, 2007 Printing Time: 9:38:57 AM

225 of 281

-129-

Operator: Shelley Hollar nalyst Version: 1.4.2 Table A-12: I6-534006l1 Data

-130-226 of 281

Note:

Experimental urine sample analysis

Study Record Page: 224b

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227 of 281

Resu	lts Path: \\Lcmsp(	03\scie	xdata\Projects\53	34006\Bio\R	esults\l6-53	34006	1.rdb											Page 1 of 2
	File Name	Sample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record	Analyte Concentration (ng/mL)	Dilution Factor	Calculated Concentration (no/mL)	%RE	Analyte Peak Area (counts)		Analyte Integration	Record Modified	Sample Annotation	LCARTRIDGET RAY	LCARTRIDGEP OSITION	
	16-53400611\16-1254.wiff		Test Solution	Unknown		<u> </u>	N/A	1.00	No intercept			5.93	Valley			10	A1	
	6-534006 1\l6-1255.wilf  6-534006 1\l6-1256,wilf		Test Solution Test Solution	Unknown		ļ	N/A N/A	1.00	No Intercept	#BAD!	2.98e+007 2.96e+007	5.93 5.94	Base To Base	1	peak splitting factor		B1 C1	
	6-53400611\l6-1257.wiff		Test Solution	Unknown		ļ	N/A	1.00	No Intercept No Intercept	#BAD!	2.96e+007	5.99	Base To Base Base To Base		ipeak spilling factor		C1	
5	6-534006I1\I6-1258.wiff		Test Solution	Unknown		ļ	N/A	1.00	No Intercept	#BAD!	2.99e+007	5.98	Base To Base		peak splitting factor		C1	
6	6-534006 1\l6-1259.wiff		Test Solution	Unknown	•	i	N/A	1.00	No Intercept	#BADI	3.02e+007	6.00	Base To Base		· [	10	Λ1	
	16-53400611\16-1260.wiff		Test Solution	Unknown		<u> </u>	N/A	1.00	No Intercept	#BAD!	3.03e+007	6.01	Base To Base				A1	
	16-53400611\16-1261.wiff 16-53400611\16-1262.wiff		Test Solution Test Solution	Unknown		ļ	N/A N/A	1.00	No Intercept	#BAD!	3.02e+007 3.05e+007	5.92 5.99	Base To Base			.1	A1	
	6-53400611\l6-1263.wiff		Test Solution	Unknown		<b></b>	N/A	1.00	No Intercept No Intercept	#BADI	3.05e+007 3.08e+007	6.00	Base To Base Base To Base	$\parallel$ $\parallel$			A1	
	6-53400611\16-1264.wiff		Mobile Phase	Unknown		ļ	N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak	<del>                                     </del>	·		D1	
12	6-534006 1\l6-1265.wilf	220-1	Solvent Blank	Unknown		······	N/A	1.00	No Peak	N∕A	0.00e+000	0.00	No Peak	1 5	·		E1	
	6-534006 1\l6-1266.wilf		Blank Urine	Unknown		<u></u>	N/A	1.00	< 0	#BAD!	5.96e+004	6.11	Base To Base			10	F1	
	16-53400611\16-1267.will		Blank Urine	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak				G1	
15 16	6-534006  1\16-1268, wilf  6-534006  1\16-1269, wilf		Blank Urine Mobile Phase	Unknown		ļ	N/A N/A	1.00	No Peak No Peak	N/A N/A		0.00	No Peak				H1	
		216-1	C 10	Standard		[X]	10.000	1.00	10.583	5.8		6.14	No Peak Base To Base				G2 C3	
		216-4	C 30	Standard		<u>     </u>	30.000	1.00	28.465	-5.1	6.93e+005	6.09	Base To Base	╁╌	·		F3	
19	16-534006(1\16-1272.wiff	216-7	C 100	Standard		<b>X</b>	100.00	1.00	100.87	0.87	2.16e+006	6.14	Valley	1 5			Н3	
20	6-534006 1\l6-1273.wilf		C 300	Standard		⊠	300.00	1.00	321.35	7.1	6.33e+006	6.16	Base To Base	⊠	peak splitting factor		C4	
	6-534006 1\16-1274.wiff		C 500	Standard		⋈	500.00	1.00	510.66	2.1	9.55e+006	6.12	Base To Base			1	C4	
22 23	6-534006  1\16-1275,wilf  6-534006  1\16-1276,wilf		C 750 C 1000	Standard Standard		× ×	750.00 1000.0	1.00	731.66 1132.4	-2.4 13.	1.29e+007 1.77e+007	6.16 6.02	Base To Base Base To Base	<u> </u>	<b></b>		C4 F4	
24	6-53400611\16-1277.will		Mobile Phase	Unknown		☒	N/A	1.00	No Peak	N/A	0.00e+000	0.02	No Peak				C6	
3	6-534006 1\16-1278.will		QC 30	Quality Control	_	×	30.000	1.00	27.867	-7.1	6.81e+005	6.11	Base To Base				E6	
26	6-534006 1\l6-1279.wilf	217-4	QC 250	Quality Control	••••	×	250.00	1.00	226.99	-9.2	4.60e+006	6.09	Base To Base	18	· · · · · · · · · · · · · · · · · · ·	10	G6	
	6-534006 1\l6-1280.wiff		QC 750	Quality Control		×	750.00	1.00	769.59	2.6		6.09	Base To Base		·		D7	
	6-534006 1\ 6-1281.wifi		QC 30000	Quality Control		☒	30000.	1000.	28222.	-5.9		6.15	Base To Base				A7	
	6-534006 1\ 6-1282.will  6-534006 1\ 6-1283.will		Mobile Phase QC 30	Unknown Quality Control			N/A 30.000	1.00	No Peak 31.175	N/A 3.9	0.00e+000 7.49e+005	0.00 6,10	No Peak			***· I	D1 E6	
	6-53400611\16-1284.wilf		QC 250	Quality Control	_	×	250.00	1.00	240.16	-3.9	4.85e+006	6.04	Base To Base Base To Base	$\vdash \vdash$			G6	
	6-534006I1\I6-1285.will		QC 750	Quality Control	****	⊠	750.00	1.00	750.46	0.062	1.31e+007	6.00	Valley	╁╌	-	1	D7	
33	6-534006(1\16-1286.will		QC 30000	Quality Control	****	Ø	30000.	1000.	27523.	-8.3	6.74e+005	6.03	Base To Base	1 7	·	10	A7	
	6-534006  1\ 6-1287.will		Mobile Phase	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			1	D1	
	6-534006I1\l6-1288.wiff		46664, 0-6 hr	Unknown			N/A	1000,	345340.	N/A	6.76e+006	6.09	Base To Base			. † '	D1	
	6-53400611\16-1289.wiff  6-53400611\16-1290.wiff		46664, 0-6 hr rinse 46667, 0-6 hr rinse	Unknown		ļ	N/A N/A	1000. 1000.	6510.1 96774.	N/A N/A	2,38e+005 2.08e+006	6.10 6.09	Base To Base Base To Base		<u> </u>		D1	
	6-534006I1\I6-1291.wiff		46670, 0-6 hr	Unknown			N/A	1000.	394510.	N/A	7.61e+006	6.09	Base To Base	<del>- H</del> -		1	D1	
	6-534006 1\16-1292.wiff		46682, 0-6 hr	Unknown	tame.		N/A	1000.	564250.	N/A	1.04e+007	6.24	Base To Base	<del>                                     </del>		10	D1	
	6-534006 1\16-1293.wiff		46683, 0-6 hr	Unknown	****		N/A	1000.	1394900.	N/A	2.01e+007	6.17	Valley		1	1.0	D1	
	6-53400611\l6-1294.wiff		46683, 0-6 hr rinse	Unknown	****		N/A	1000.	12178.	N/A	3.56e+005	6.15	Base To Base				D1	
42 43	6-53400611\l6-1295,wiff 6-53400611\l6-1296.wiff		46690, 0-6 hr 46664, 6-12 hr	Unknown			N/A N/A	1000.	394380. 141640.	N/A N/A	7.61e+006 2.97e+006	6.14 6.16	Base To Base			.1	D1	
	6-53400611\6-1297.wiff		46667, 6-12 hr	Unknown	*****	ļ	N/A	1000.	104600,	N/A	2.976+006 2.24e+006	6.22	Base To Base Base To Base			1	D1	
	6-53400611\l6-1298.wiff		46670, 6-12 hr	Unknown	****	<b></b>	N/A	1000.	89395.	N/A	1.93e+006	6.17	Base To Base				D1	
	6-534006 1\l6-1299.wiff		46682, 6-12 hr	Unknown		<b></b>	N/A	1000.	6970.5	N/A	2.48e+005	6.12	Base To Base	╁┼	<del> </del>	10	D1	
	6-534006  1\16-1300,wiff		46683, 6-12 hr	Unknown			N⁄A	1000.	74686.	N/A	1.64e+006	6.04	Base To Base				D1	
	6-534006(1\l6-1301,wiff		46690, 6-12 hr	Unknown			N/A	1000.	113540.	N/A	2.41e+006	6.17	Base To Base				D1	
	6-53400611\l6-1302.wiff 6-53400611\l6-1303.wiff		46664, 12-24 hr 46667, 12-24 hr	Unknown Unknown	*****		N/A N/A	1000. 1000.	8081.7 15157.	N/A N/A	2.71e+005 4.18e+005	6.11	Base To Base				D1	
	6-53400611\6-1303,Will		46670, 12-24 hr	Unknown	••••	ļ	N/A N/A	1000.	8781.7	N/A	4.18e+005 2.85e+005	6.20 6.22	Base To Base Base To Base	<del>                                     </del>	<b> </b>		D1	
	6-534006l1\16-1305.wiff		Mobile Phase	Unknown		ļ	N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak	<del>- H</del> -	<del> </del>		D1	
53	6-534006  1\16-1306.wiff	216-2	C 10	Standard		☒	10.000	1.00	11.390	14.	3.40e+005	6.23	Base To Base	H	<del> </del>	1	E3	
	G-534006 1\l6-1307.wilf		C 30	Standard		⊠	30.000	1.00	29.241	-2.5		6.22	Base To Base		†		F3	
	6-534006 1\ 6-1308.wi		C 100	Standard	*****	⊠	100.00	1.00		3.1		6.26	Base To Base				B4	
	6-53400611\l6-1309.wilf		C 300	Standard	*****	×	300,00			2.2		6.22	Base To Base		ļ		E4	
57 58	6-53400611\\6-1310.wiff 6-53400611\\6-1311.wiff		C 500 C 750	Standard Standard		<u> </u>	500.00 750.00	1.00		-5.5 0.46		6.17 6.15	Base To Base Base To Base				E4	
0.003 (33)	6-53400611\(6-1312.will		C 1000	Standard		⊠ ⊠	1000.0	1.00	1014.0	1.4		6.18	Valley	<del>                                     </del>	<b>}</b>	1	G4	
	6-534006I1\l6-1313.wiff		Mobile Phase	Unknown	****	<u> </u>	N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak	╀┈	<del> </del>		C6	
61	6-534006 1\ 6-1314.wiff	217-2	QC 30	Quality Control		՛՛⊠	30.000	1.00	26.876	-10.	6.60e+005	6.23	Base To Base		<del> </del>	1	E6	
62	6-534006 1\l6-1315.wiff		QC 250	Quality Control		Ø	250.00	1.00	207.90	A	4.24e+006	6.23	Base To Base			1'I	G6	
63	6-53400611\l6-1316.wiff		QC 750	Quality Control		☒	750.00	1.00	758.41	1.1		6.22	Base To Base			1.0	D7	
	6-53400611\l6-1317.wiff 6-53400611\l6-1318.wiff		QC 30000 Mobile Phase	Quality Control Unknown		☒	30000.	1000.	25817.			6.22	Base To Base	⊠	peak splitting factor	1	A7	
	6-53400611\16-1319.will		QC 30	Quality Control		Ø	N/A 30.000	1.00	No Peak 29.406			0.00 6.15	No Peak Base To Base		ļ	1	D1 E6	
الانتشاء			<del></del>	1				L	L	1		12.12	1200 10 0000		L	<u></u>		

	File Name	Sample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record	Analyte Concentration (ng/mL)	Dilution Factor	Calculated Concentration (no/mL)	%RE	Analyte Peak Area (counts)	Analyte Retention Time (min)	Analyte Integration Type	Record Modified	Sample Annotation	LCARTRIDGET RAY	LCARTRIDGEP OSITION
	16-53400611\16-1320,\viff		QC 250	Quality Control	<u> </u>	⊠	250.00	1.00	232,17	-7.1		6.09	Base To Base	П	<u> </u>	10	IG6
	16-53400611\16-1321.will		QC 750	Quality Control	1	×	750.00	1.00	783.18	4.4	1.36e+007	6.14	Base To Base			10	D7
	16-534006  1\16-1322.wiff		QC 30000	Quality Control		⊠	30000.	1000.	28519.	-4.9	6.94e+005	6.07	Base To Base		***************************************	10	A7
	16-534006l1\16-1323.wiff	·	Mobile Phase	Unknown	1		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			10	D1
	16-53400611\16-1324.wiff	I	46670, 0-6 hr rinse	Unknown	1		N/A	100.	4827.7	N/A	1.10e+006	6.21	Base To Base			10	D1
	16-53400611\16-1325.wiff		46682, 0-6 hr rinse	Unknown	1		N/A	100.	8214.3	N/A	1.79e+006	6.17	Base To Base			10	D1
	16-534006  1\16-1326, will		46690, 0-6 hr rinse	Unknown	1 -		N/A	1D0.	7204.7	N/A	1.58e+006	6.24	Base To Base	i i	······································	10	D1
	l6-534006l1\l6-1327.wiff		46664, 6-12 hr rinse	Unknown	1		N/A	100.	3614.2	NA	8.51e+005	6.20	Base To Base	П		10	D1
	16-534006 1\16-1328.will		46667, 6-12 hr rinse	Unknown	· · · ·	~~~~~	N/A	100.	1849.0	N/A	4.87e+005	6.22	Base To Base			10	D1
	16-53400611\16-1329.wiff		46670, 6-12 hr rinse	Unknown	1 -		N/A	100.	2294.6	NA	5.79e+005	6.09	Base To Base	n i		10	D1
	16-53400611\16-1330,wiff		46683, 6-12 hr rinse	Unknown	·		N/A	100.	2117.5	N/A	5.43e+005	6.22	Base To Base			10	D1
	16-53400611\16-1331.wiff		46690, 6-12 hr rinse	Unknown			N/A	100.	2037.8	N∕A	5.26e+005	6.25	Base To Base			10	D1
	16-534006 1\16-1332.will		46667, 12-24 hr rinse	Unknown	1		N/A	100.	948.30	N/A	3.00e+005	6.23	Base To Base	i i	***************************************	10	D1
	l6-534006 1\l6-1333.wilf		46682, 12-24 hr	Unknown			N/A	100.	669.74	N/A	2.42e+005	6.20	Base To Base	- H		10	D1
	16-534006  1\16-1334.wiff		46683, 12-24 hr	Unknown			N/A	100.	767.00	N/A	2.62e+005	6.20	Base To Base		***************************************	10	D1
	16-534006  1\16-1335.wiff		Mobile Phase	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			10	C6
	16-534006 1\16-1336.wiff		QC 30	Quality Control		⊠	30.000	1.00	24.937	-17.	6.20e+005	6.28	Base To Base	i i		10	E6
34	16-53400611\16-1337.wiff	217-6	QC 250	Quality Control		⊠	250.00	1.00	228.59	-8.6	4.63e+006	6.30	Base To Base		·	10	G6
	16-534006(1\16-1338,wiff		QC 750	Quality Control		⊠	750.00	1.00	745.96	-0.54	1.31e+007	6.27	Base To Base			10	D7
للشششية	16-534006  1\16-1339.wiff	L	QC 30000	Quality Control		×	30000.	1000.	26170.	-13.	6.46e+005	6.24	Base To Base			10	A7
	16-53400611\16-1340.wiff		Mobile Phase	Unknown	-		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak	- Fi		10	D1
18	16-53400611\16-1341.wiff	219-3	QC 30	Quality Control		⊠	30.000	1.00	29.927	-0.24	7.23e+005	6,14	Base To Base	l fil	······	10	E6
19	16-534006  1\16-1342.wiff	219-6	QC 250	Quality Control		<b>(X)</b>	250.00	1.00	252.99	1.2	5.09e+006	6.12	Base To Base		······································	10	G6
	16-534006l1\16-1343.wiff		QC 750	Quality Control		X	750.00	1.00	783,15	4.4	1.36e+007	6.14	Base To Base	h ii		10	D7
	16-534006  1\16-1344.viif		QC 30000	Quality Control		×	30000.	1000.	27956.	-6,8	6.83e+005	6.10	Base To Base	H H		10	A7
2	16-53400611\16-1345.wiff	213-1	Mobile Phase	Unknown		,	N/A	1,00	No Peak	N/A	0.00e+000	0.00	No Peak	H		10	D1
3	16-53400611\16-1346.wiff	216-3	C 10	Standard	*****	⊠	10.000	1.00	8.0845	-19.	2.71e+005	6.28	Base To Base		peak splitting factor	10	F3
4	16-53400611\16-1347.wiff	216-6	C 30	Standard		×	30.000	1.00	31.452	4.8	7.55e+005	6.21	Base To Base	H H		10	F3
15	16-53400611\16-1348.wiff	216-9	C 100	Standard		⊠	100.00	1.00	99.180	-0.82	2.13e+006	6.31	Base To Base	H		10	B4
6	IG-534006l1\I6-1349.wiff	216-12	C 300	Slandard		⊠	300,00	1.00	299.54	-0.15	5.94e+006	6.21	Base To Base		peak splitting factor	10	E4
	16-534006  1\16-1350.wiff		C 500	Standard			500.00	1.00	459.37	-8.1	8.71e+006	6.27	Base To Base	ñ			E4
8	16-534006  1\16-1351.wiff	216-18	C 750	Standard		$\boxtimes$	750.00	1.00	692.61	-7.7	1.23e+007	6.27	Base To Base		~~~~~~~~		E4
	16-534006  1\16-1352.wiff		C 1000	Standard		☒	1000.0	1.00	984.27	-1.6	1.61e+007	6,29	Base To Base	- H		10	G4
00	16-53400611\16-1353.will	213-1	Mobile Phase	Unknown	·		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak	H	***************************************		D1

-133-229 of 281

> Printing Date: Friday, March 30, 2007 Printing Time: 8:07:23 AM

Operator: Shelley Hollar nalyst Version: 1.4.2

-15<del>4-</del> 230 of 281

Note:

Experimental urine sample analysis

Freeze-thaw cycle, 14-day frozen and 4 hour benchtop stability assessments.

534006-227-4 was not included in the regression calculations based on the test for outliers.

Study Record Page: 237a

			T	T				·					(0.000.000.000.000.000.000.000.000.000.		100000000000000000000000000000000000000
	File Name	Sample	Sample Name	Sample Type	Analyte Peak	U5e	Analyte Concentration	Dilution	Calculated Concentration	%RE	Analyte Peak	Analyte Retention	Analyte Integration	Record	Sample Annotation
1,20	170 L. S. S. S. S. S. S. S. S. S. S. S. S. S.	, ID	Lample Name	Sample 1996	Name :	Record	(ng/mL)	Factor	(rig/mL)	A047.28	Area (counts)	Time (min)	Тура	Modified	Janipie Amotation
	16-534006m\16-1354,wiff	200220777	Test Solution	Unknown	Window	MAX	NA	1.00	No Intercept	#BADI	1.73e+007	6.09	Base To Base		De 1000000000000000000000000000000000000
•	16-534006m\16-1355.wiff		Test Solution	Unknown	-		N/A	1.00	No Intercept	#BADI	1.6Be+007	6.00	Base To Base	<del>                                     </del>	ļ
3	16-534006m\16-1356.vvilf		Test Solution	Unknown	-		N/A	1.00	No Intercept	#BAD!	1.70e+007	5.97	Base To Base	╁┼	ļ
4	16-534006m\16-1357,wiff		Test Solution	Unknown			N/A	1.00	No Intercept	#BAD!		6.01	Base To Base		ļ
	16-534006m\16-1358.wiff	ļ	Test Solution	Unknown	-		N/A	1.00	No intercept	#BAD!		5.96			<b></b>
9					-								Base To Base	<u> </u>	<u> </u>
6	16-534006m\16-1359.wiff		Test Solution	Unknown			N/A	1.00	No Intercept	#BAD!		6.04	Base To Base		
7	16-534006m\16-1360.wiff		Test Solution	Unknown			N/A	1.00	No Intercept	#BAD!		5.95	Base To Base	⊠	peak splitting factor
8	l6-534006m\l6-1361.wilf		Test Solution	Unknown	_		N/A	1.00	No intercept	#BAD!	1.80e+007	5.95	Base To Base		
9	16-534006m\16-1362.wiff		Test Solution	Unknown	_		N/A	1.00	No intercept	#BAD!		5.97	Base To Base		
10	16-534006m\16-1363.wiff		Test Solution	Unknown			N/A	1.00	No Intercept	#BAD!		5.93	Base To Base		
11	l6-534006m\l6-1364.wiff		Mobile Phase	Unknown			N/A	1.00	No Peak	NVA	0.00e+000	0.00	No Peak		
12	16-534006m\16-1365.wiff		Solvent Blank	Unknown	_			1.00	No Peak	NVA		0.00	No Peak		<u> </u>
13	16-534006m\16-1366.wiff		Blank Urine	Unknown			N/A	1.00	No Peak	NA		0,00	No Peak		l
14			Blank Urine	Unknown					No Peak	N/A		0.00	No Peak		
15			Blank Urine	Unknown	l ï		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		1
16			Mobile Phase	Unknown			N/A		No Peak	N/A		0.00	No Peak		T
17	l6-534006m\l6-1370.wiff	227-1	C 10	Standard		⊠	10.000	1.00	9.2867	-7.1	9.66e+004	5.82	Base To Base		T
18	16-534006m\16-1371,wifl		C 30	Standard			30.000		67.890	130.	6.792+005	5.77	Base To Base		<b>1</b>
19	16-534006m\16-1372,wiff	227-7	C 100	Standard	-	Ø	100.00	1.00	87.750	-12.	8.74e+005	5.88	Base To Base		1 .
20	16-534006m\16-1373,wiff	227-10	C 300	Standard	-	×	300.00	1.00	311.93	4.0	2.95e+006	5.90	Base To Base		
21	16-534006m\16-1374.wiff	227-13	C 500	Standard	-	⊠	500.00	1.00	516,33	3.3	4.65e+006	5.88	Base To Base	1 7	<u> </u>
22	16-534006m\l6-1375.wiff	227-16	C 750	Standard	-		750.00	1.00	757.42	0.99	6.42e+006	5.88	Base To Base		·
23	16-534006m\16-1376.wiff	227-19	C 1000	Standard	-		1000.0	1.00	918.88	-8.1	7,47e+006	5.86	Base To Base		
24	16-534006m\l6-1377.wiff	234-1	Mobile Phase	Unknown	-		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak	1 5	ļ
25	16-534006m\16-1378.wiff	228-1	QC 30	Quality Control	-	⊠	30.000	1.00	28.834	-3.9	2.93e+005	5.91	Base To Base	1-5-	İ
26			QC 250	Quality Control	-		250.00		224,67			5.90	Valley		<del> </del>
27			QC 750	Quality Control	-				645.14	-14.		5.78	Base To Base	<del>                                     </del>	ļ
28			QC 150000	Quality Control	-			5000.	173620.	16.		5.82	Base To Base	1 8	<del> </del>
29			Mobile Phase	Unknown	-		N/A		No Peak	N/A		0.00	No Peak	┨	ļ
30			46683, 0-6 hr	Unknown	-		N/A	5000.	1533800.	N/A		5.87	Base To Base		
31			Mobile Phase	Unknown	-		N/A		No Peak	N/A	0.002+000	0.00	No Peak		
32			46664, 0-6 hr rinse	Unknown			N/A		10363.	N/A		5.85	Base To Base	<u> </u>	
33			46682, 6-12 hr	Unknown	-		N/A		683.33	N/A		5.74	Base To Base	<u> </u>	
34			Mobile Phase				N/A							<u>                                     </u>	
35			46682, 12-24 hr	Unknown			N/A		No Peak	N/A #BAD!		0.00	No Peak		ļ
100				Unknown					No Intercept			5.89	Base To Base	<u> </u>	
36			46683, 12-24 hr	Unknown			N/A	2.00	1315.4	N/A		5.92	Base To Base	↓ □	ļ
37			Mobile Phase	Unknown			N/A	1.00	No Peak	N/A		0.00	No Peak		<b></b>
38			30 ng/mL 4 Hr Rt Stb	Unknown			N/A		31.238	N/A		5.96	Base To Base	<u> </u>	<b></b>
39			30 ng/mL 4 Hr Rt Slb	Unknown	_		N/A		28.626	N/A	2.91e+005	5.89	Base To Base		
40			30 ng/mL 4 Hr Rt Stb	Unknown	_		N/A		26.302	N/A	2.87e+005	5.93	Base To Base		<u> </u>
41			750 ng/mL 4 Hr Rt Stb	Unknown			N/A	1,00	761.39	N/A	6.45e+006	5.88	Base To Base	⊠	peak splitting factor
42			750 ng/mL 4 Hr Rt Slb	Unknown			N/A	1.00	765.74	N/A	6.48e+006	5.91	Base To Base		
43			750 ng/mL 4 Hr Rt Stb	Unknown	_		N/A		826.46	N/A	6.68e+006	5.89	Base To Base		<u> </u>
44			Mobile Phase	Unknown	-		N/A		No Peak	N/A	0.00e+000	0.00	No Peak		1
45	16-534006m\16-1398.wiff	227-2	C 10	Standard	_	⊠	10.000	1.00	9.5620	-4.4	9.94e+004	5.88	Base To Base		1
46	16-534006m\16-1399.wiff	227-5	C 30	Standard			30.000	1.00	28.830	-3.9	2.93e+005	6.00	Base To Base		I
47	16-534006m\l6-1400.will	227-8	C 100	Standard	-		100.00	1.00	105.56	5.6	1.05e+006	5.82	Base To Base		<u> </u>
48			C 300	Standard	-				294,75	-1.8		5.88	Base To Base		<del> </del>
49			C 500	Standard					535.69	7.1	4.80e+006	5.86	Valley		†
50			C 750	Standard	-				672.67	-10.	5.83e+006	5.84	Valley	<del>                                     </del>	<del> </del>
51			C 1000	Standard	-		1000.0	1.00	1080.6	8.1	8.40e+006	5.90	Base To Base	╁	
52		234-1	Mobile Phase	Unknown				1.00	No Peak	N/A	0.00e+000	0.00	No Peak	<del>      -   -   -   -   -   -   -   -   -</del>	
53			QC 30	Quality Control					28.265			5.89	Base To Base	├─┼┈	
				Jacany Control			·····	,.00	20.200	-5.0	2.0767003	0.03	Dose IO Dase		<u> </u>

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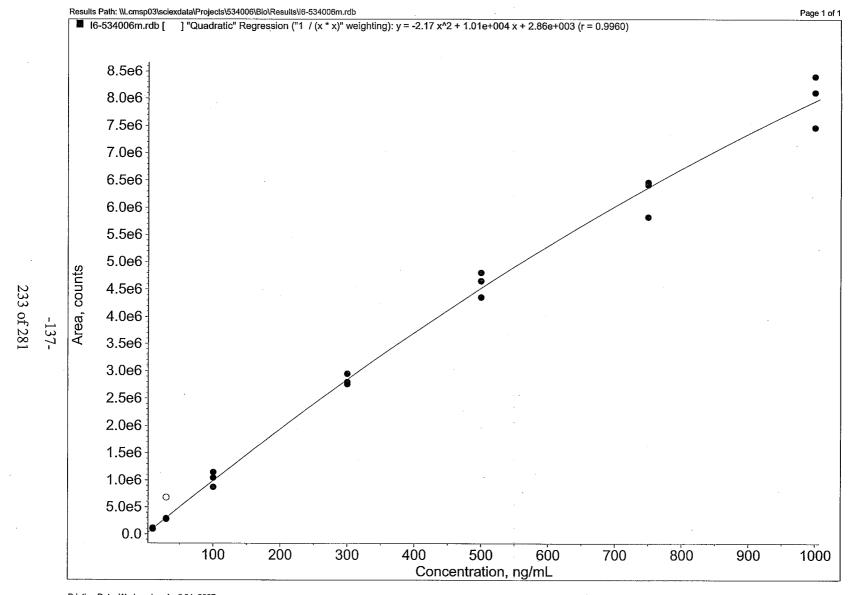
Operator: Shelley Hollar nalyst Version: 1.4.2

	File Name	Sample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record		Dilution Factor	Calculated Concentration (rig/mL)	%RE	Analyte Peak Area (counts)	Analyte Retention Time (min)	Analyte Integration Type	Record Modified	Sample Annotation
54	16-534006m\l6-1407.viff	228-5	QC 250	Quality Control		⊠	250.00	1.00	207.20	-17.	2.01e+006	5.89	Base To Base	П	
55	16-534006m\16-1408,wiff	228-8	QC 750	Quality Control		⊠	750.00	1.00	683.29	-8.9	5.90e+006	5.87	Base To Base		
56	16-534006m\16-1409.wiff	228-11	QC 150000	Quality Control	1	Ø	150000.	5000.	164740.	9.8	3.34e+005	5.86	Base To Base		peak splitting factor
57	16-534006m\l6-1410,wiff	234-1	Mobile Phase	Unknown	1		NA	1.00	No Peak	N/A	0,00e+000	0.00	No Peak	- H	i
58	16-534006m\16-1411.wiff	235-1	30 ng/mL F Thaw	Unknown	_		N/A	1.00	31,558	₩A	3.20e+005	5.92	Base To Base	Ħ	
59	16-534006m\16-1412.wiff	235-2	30 ng/mL F Thaw	Unknown			N/A	1.00	30.036	N/A	3.05e+005	5.89	Base To Base		
60			30 ng/mL F Thaw	Unknown			N/A	1.00	29.277	N/A	2.97e+005	6.00	Base To Base		
61			30 ng/mL F Thaw	Unknown	•	•••••	N/A	1.00	31.244	N/A	3.17e+005	5.94	Base To Base		
62			30 ng/mL F Thaw	Unknown	7		WA	1,00	28.084	N/A	2.85e+005	5.98	Base To Base		
	16-534006m\16-1416.wiff		30 ng/ml. F Thaw	Unknown			N/A	1.00	29.979	N/A	3.04e+005	5.95	Base To Base		İ
64			30 ng/ml, F Thaw	Unknown	T		N/A	1.00	30.692	N/A	3.11e+005	5.90	Base To Base		İ
65	16-534006m\16-1418.wiff		30 ng/mL F Thaw	Unknown	_		N/A	1.00	28.488	N/A	2.89e+005	5.99	Base To Base		
66	16-534006m\16-1419.wiff	235-9	30 ng/mL, F Thaw	Unknown			N/A	1.00	29.403	N/A	2.98e+005	5.95	Base To Base		
67	16-534006m\16-1420.wiff	235-10	750 ng/mL, F Thaw	Unknown	"		N/A	1.00	808.71	N/A	6.77e+006	5.96	Base To Base	П	
68	16-534006m\16-1421.wiff		750 ng/mL F Thaw	Unknown	,		WA	1.00	878.47	N/A	7.22e+006	5.95	Base To Base		
69	16-534006m\16-1422.wiff	235-12	750 ng/mL F Thaw	Unknown			N/A	1.00	830.72	N/A	6.91e+006	5.99	Base To Base		
70	16-534006m\16-1423.wiff		750 ng/mL F Thaw	Unknown	-		N/A	1.00	861.41	ΝA	7.11e+006	5.95	Base To Base		······································
71 ::	16-534006m\16-1424.wiff	235-14	750 ng/mL F Thaw	Unknown	-		N/A	1.00	899.14	N/A	7.35e+006	5.95	Base To Base		
72	16-534006m\16-1425.wiff	235-15	750 ng/mL F Thaw	Unknown	·		N/A	1.00	668.23	N/A	5.79e+006	5.96	Base To Base		
73	16-534006m\16-1426.will	235-16	750 ng/mL F Thaw	Unknown			N/A	1.00	822.95	N/A	6.86e+006	5.95	Base To Base		
74	16-534006m\16-1427.will		750 ng/mL F Thaw	Unknown	_		N/A	1.00	759.15	N/A	6.43e+006	5.68	Base To Base		
75	16-534006m\l6-1428.wiff		750 ng/mL F Thaw	Unknown	-		N/A	1.00	870.17	NVA	7.16e+006	5.97	Base To Base		
76	16-534006m\16-1429.wiff		Mobile Phase	Unknown	"		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak	n	<del></del>
77	16-534006m\l6-1430.wiff		QC 30	Quality Control	_	⊠	30.000	1.00	27.496	-8.3	2.79e+005	5.92	Base To Base		i
78	16-534006m\16-1431.wiff	228-6	QC 250	Quality Control		⊠	250.00	1.00	265.19	6.1	2.53e+006	5.85	Base To Base		
79	l6-534006m\l6-1432.wiff		QC 750	Quality Control	٦	(⊠	750,00	1.00	816.12	9.1	6.83e+006	5.85	Base To Base		
80	l6-534006m\l6-1433.wiff	228-12	QC 150000	Quality Control	٦ ٦	⊠	150000.	5000.	160180.	6.8	3.25e+005	5.89	Base To Base	<u>×</u>	peak splitting factor
81	16-534006m\16-1434.wiff			Unknown	1		N/A		No Peak	NΑ	0.00e+000	0.00	No Peak		
82	16-534006m\l6-1435.wiff			Standard	7	⊠	10.000	1.00	11.459	15.	1.18e+005	5,91	Valley		
83				Standard	- 1	⊠	30.000	1.00	27.572	-8.1	2.80e+005	5.87	Base To Base		
84	l6-534006m\l6-1437.wiff			Standard			100.00	1.00	116.01	16.	1.15e+006	5.94	Base To Base		
85	l6-534006m\l6-1438.wiff			Standard	7		300.00	1.00	290,51	-3.2	2.76e+006	5.94	Base To Base		
86	16-534006m\16-1439.wiff			Standard		⊠	500.00	1.00	479,39	-4.1	4.35e+006	5.91	Base To Base		
87	16-534006m\l6-1440.wiff		C 750	Standard	7	⊠	750.00	1.00	763,54	1.8	6.46e+006	5.94	Base To Base		
	16-534006m\16-1441.wiff		C 1000	Slandard	7	×	1000.0	1.00	1027.2	2.7	8.11e+006	6.00	Base To Base		
69	16-534006m\16-1442.wiff	234-1	Mobile Phase	Unknown	7		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		

Printing Date: Wednesday, April 04, 2007 Printing Time: 8:26:26 AM

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Printing Date: Wednesday, April 04, 2007

Operator: Shelley Hollar

-138-234 of 281

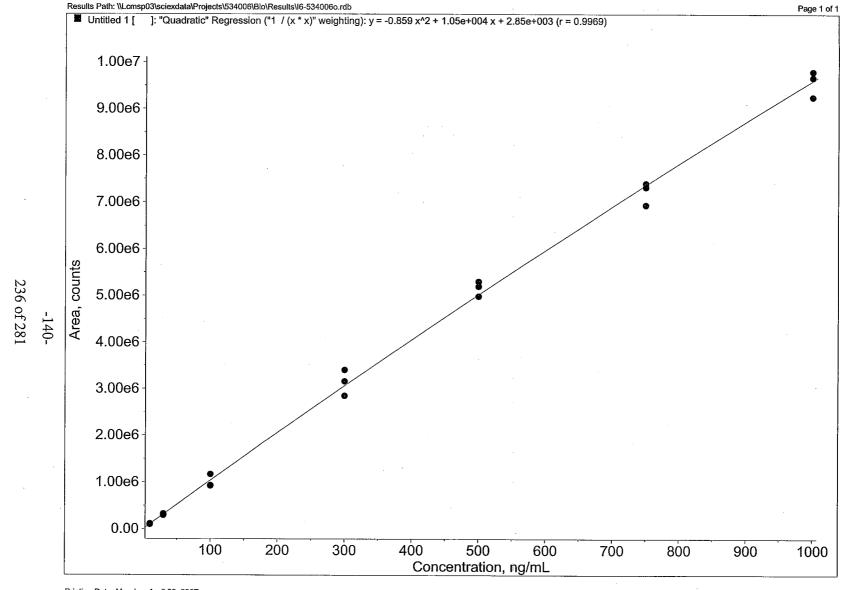
Note

Experimental urine sample analysis

Study Record Page: 256a

	File Name	Sample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record	Analyte Concentration (ng/mL)	Dilution Factor	Calculated Concentration (ng/mL)	%RE	Analyte Peak Area (counts)	Analyte Retention Time (min)	Analyte Integration Type	Record Modified	Sample Annotation
. X	16-534006o\16-1514.wiff	248-10	Test Solution	Unknown		20020000	N/A	1.00	1696.2	WA	1.53e+007	5.78	Base To Base		
	16-534006o\16-1515.wiff	248-10	Test Solution	Unknown		<b></b>	N/A	1.00	1795.2	N/A	1.60e+007	5.83	Base To Base	-	
		<u></u>	Test Solution	Unknown			N/A	1.00	1799.3	N/A	1.61e+007	5.69	Base To Base		
		248-10	Test Solution	Unknown			N/A	1.00	1872.9	N/A	1.66e+007	5.74	Base To Base	<u> </u>	
	16-534006o\16-1518.wiff	L	Test Solution	Unknown	·		N/A	1.00	1844.3	N/A	1.64e+007	5.75	<u> </u>	<u> </u>	
120.0			Test Solution	Unknown			N/A	1.00	1934.7	N/A			Base To Base	<u> </u>	
		248-10	Test Solution	Unknown			N/A	1.00	1956.5	N/A	1.70e+007	5.76	Base To Base	<u> </u>	
			Test Solution	Unknown			N/A	1.00	1947.1			5.72	Base To Base		ļ
		248-10	Test Solution	Unknown			N/A	1.00		N/A	1.71e+007	5.72	Base To Base		<u></u>
	16-5340060\16-1523.wiff	248-10	Test Solution						2016.4	N/A	1.76e+007	5.68	Base To Base		
			I	Unknown			N/A	1.00	2169.6	N/A	1.87e+007	5.68	Base To Base		
75,71		256-1	Mobile Phase	Unknown	********		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
	I6-534006o\I6-1525,wiff	252-1	Solvent Blank	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
		252-2	Blank Urine	Unknown	*******		N/A		No Peak	N/A	0.00e+000	0.00	No Peak		
		252-3	Blank Urine	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
		252-4	Blank Urine	Unknown			N/A	1.00	No Peak	N/A		0.00	No Peak		
			Mobile Phase	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
			C 10	Standard		⊠	10.000	1.00	9.1005	-9.0	.i	5.54	Base To Base	⊠	noise % peak splilling
			C 10	Standard		×	10.000	1.00	10.345	3.5	1.11e+005	5.62	Base To Base	⊠	noise % peak splitting
			C 30	Standard	******	☒	30.000	1.00	31.185	4.0	3.28e+005	5.62	Base To Base	⊠	peak splitting ratio
			C 30	Standard		☒	30.000	1.00	27.949	-6.8	2.95e+005	5.60	Base To Base		
			C 100	Standard		⊠	100.00	1.00	89.411	-11.	9.32e+005	5.52	Base To Base		
ķ.	16-534006o\16-1535.will	250-8	C 100	Slandard		⊠	100.00	1.00	88.591	-11.	9.23e+005	5.54	Base To Base		
	16-534006o\16-1536.will	250-10	C 300	Standard		⊠	300.00	1.00	278.52	-7.2	2.85e+006	5.60	Base To Base		***************************************
	16-534006o\16-1537.wiff	250-13	C 500	Standard	,,,,,,,	⊠	500.00	1.00	496.08	-0.78	4.98e+006	5.58	Base To Base		
	16-534006o\16-1538.wiff	250-16	C 750	Standard		⊠	750.00	1.00	703.34	-6.2	6.94e+006	5.52	Base To Base		
	16-534006o\16-1539.wiff	250-19	C 1000	Standard		⊠	1000.0	1.00	959.94	-4.0	9.26e+006	5.53	Base To Base		
	16-534006o\16-1540.wiff	256-1	Mobile Phase	Unknown	*******	***************************************	N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
	16-534006o\16-1541.wiff	251-1	QC 30	Quality Control	******	⊠	30.000	1.00	30.061	0.20	3.17e+005	5.57	Base To Base		
	16-534006o\16-1542.wiff	251-2	QC 30	Quality Control		×	30.000	1.00	26.901	-10,	2.84e+005	5.64	Base To Base		peak splitting factor
	16-534006o\16-1543.wiff	251-4	QC 250	Quality Control		Ø	250.00		252.84	1.1		5.55	Base To Base		part opining laster
-	16-534006o\16-1544.wiff	251-5	QC 250	Quality Control	*******	<u> </u>	250.00		258.15	3.3	2.65e+006	5.54	Base To Base		
T	16-534006o\16-1545.wiff		QC 750	Quality Control		Ø	750.00		807.06	7.6			Base To Base		
	16-534006o\16-1546.wiff	251-10	QC 30000	Quality Control	Material	Ø	30000.		29442.	-1.9			Base To Base		
			Mobile Phase	Unknown	******		N/A	l	No Peak	N/A		0.00	No Peak	<del>                                     </del>	
			46682, 6-12 hr	Unknown			N/A	<b></b>	12069.	N/A	J	5.52	Base To Base		
			46682, 12-24 hr	Unknown					1200.3	N/A		5.45	Base To Base		
			Mobile Phase	Unknown	,,,,,,,		N/A		No Peak	N/A		0.00	No Peak	<u> </u>	·····
			QC 30	Quality Control		⊠	30.000		27.302	-9.0		5.53	Base To Base	<u> </u>	
			QC 250	Quality Control		<u> </u>	250.00		224.11	-10.		5.53	Base To Base	Ł	
			QC 750	Quality Control	••••	⊠	750.00	1.00	735,91	-10. -1.9		5.61	Base To Base		
			QC 750	Quality Control		⊠	750.00		731.69	-2.4		5.60	Base To Base		
			QC 30000	Quality Control		_ <u>W</u> _	30000.		28811.	-4.0		5.54	Base To Base	<u></u>	····
			QC 30000	Quality Control		⊠ ⊠	30000.		32312.	7.7		5.47	Base To Base		
			Mobile Phase	Unknown			N/A		No Peak	N/A		0.00	No Peak		
			C 10	Standard		674	10.000	1.00	10.604		1		***************************************		
	16-5340060\l6-1559.wilf		C 30	.)	******	<u> </u>				6.0			Base To Base		
			C 100	Standard		<u> </u>	30.000		30,970	3.2			Base To Base		
				Standard		<u> </u>	100.00	1,00	112.69	13,			Base To Base		
			C 300	Standard	<b></b> .	⊠			334.20	11.			Base To Base		·
			C 300	Standard		⊠	300.00		309.52	3.2			Base To Base		
			C 500	Standard	,	⊠			528.93	5.8	1		Base To Base		
			C 500	Slandard					518.22	3.6	[		Base To Base		
			C 750	Standard					753.65	0.49	7.40e+006	5.55	Base To Base		
	16-534006a\16-1566.wiff		C 750	Standard		⊠			744.89	-0.68			Base To Base ·		
			C 1000	Standard	-	×	1000.0		1007.4	0.74	9.67e+006	5.50	Base To Base		
			C 1000	Standard	<del></del>	☒	1000.0	1.00	1021.9	2.2	9,80e+006		Base To Base		
100	16-534006o\16-1569.wiff	256-1	Mobile Phase	Unknown			N/A	1.00	No Peak	N/A			No Peak		······





-141-237 of 281

Note:

- Serum Long Term Stability
- Serum Freeze Thaw Stability

Study Record Page: 266b to 266c

238	
of 281	-142-

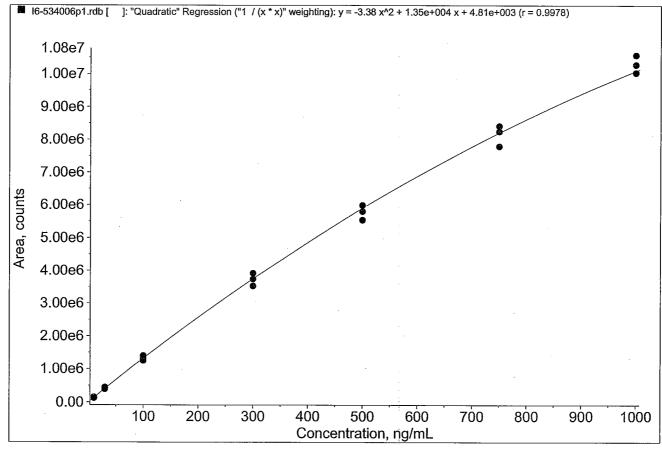
Resu	ilts Path: \\Lcmsp03\	sciexda	ata\Projects\534006	\Bio\Results\l6	5-534006p1	.rdb										Page 1 of
	File Name	Sample	Sample Name	Sample Type	Analyte Peak		Analyte Concentration	Dilution	Calculated Concentration	WOF	Analyte Peak	Analyte Retention	Analylo	Record		
	riie Name	ID	Sample Name	Sample Type	Name	Record	(ng/mL)	Factor	(ng/mL)	%RE	Area (counts)	Time (min)	Integration Type	Modified	Sample Annotation	
1	I6-534006p\I6-1647.wiff	259-10	Test Solution	Unknown	l mar	ingaeegr.nc	N/A	1.00	No Intercept	#BAD!	1.69e+007	4.66	Valley		X	
2		259-10	Test Solution	Unknown	i	İ	N/A	1.00	No Intercept	#BAD!	1.67e+007	4.63	Base To Base	177	<i></i>	
3	l6-534006p\l6-1649.wilf	259-10	Test Solution	Unknown		<u> </u>	N/A	1.00	No Intercept	#BAD!	1.72e+007	4.66	Base To Base			
4		259-10	Test Solution	Unknown			N/A	1.00	No Intercept	#BAD!	1.70e+007	4.63	Valley			
5		259-10 259-10	Test Solution	Unknown	·	ļ	N/A	1.00	No Intercept	#BAD!	1.75e+007	4.66	Base To Base			
7	16-534006p\16-1652.wiff 16-534006p\16-1653.wiff	259-10	Test Solution Test Solution	Unknown Unknown		ļ	N/A N/A	1.00	No Intercept	#BAD!	1.73e+007	4.70	Base To Base			
8		259-10	Test Solution	Unknown		<del> </del>	N/A	1.00	No Intercept No Intercept	#BADI #BADI	1.72e+007 1.70e+007	4.69 4.72	Base To Base Base To Base			
9		259-10	Test Solution	Unknown		<b>}</b>	N/A	1.00	No Intercept	#BAD!	1.68e+007	4.67	Valley			
10	i	259-10	Test Solution	Unknown		<del> </del>	N/A	1.00	No Intercept	#BADI	1.71e+007	4.71	Base To Base	╁		
11	l6-534006p\l6-1657.wiff	257-1	Mobile Phase	Unknown	- Innexes	l	N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak	╁┼┼		
12		263-1	Solvent Blank	Unknown	-	l	N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			
13	4	263-2	Blank Serum	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			
14		263-3	Blank Serum	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			
15		263-4	Blank Serum	Unknown		ļ	N/A	1.00	No Peak	N/A	0.00e+000	0,00	No Peak			
16 17		257-1	Mobile Phase	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			
18		261-1 261-4	C 10 C 30	Standard Standard		×	10,000	1.00	8.8149	-12.	1.23e+005	4.56	Valley			
19		261-4 261-7	C 100	Standard		X	30.000 100.00	1.00	28.859 94.321	-3.8 -5,7	3.91e+005 1.25e+006	4.61 4.55	Base To Base		·····	
20			C 300	Standard		X X	300.00	1.00	280.70	-5. <i>1</i> -6.4	1.25e+006 3.52e+006	4.55	Base To Base Base To Base			
21		261-13	C 500	Slandard		Ø	500.00	1.00	464.67	-7.1	5.54e+006	4.57	Base To Base		Peak splitting factor	
22			C 750	Standard	******	Ø	750.00	1.00	701.21	-6.5	7.79e+006	4.57	Base To Base		r ear aplitting factor	
23		261-19	C 1000	Standard		Ø	1000.0	1.00	991.67	-0.83	1.00e+007	4.56	Base To Base			
24	16-534006p\16-1670.wiff	257-1	Mobile Phase	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			
25		262-1	QC 30	Quality Control		⊠	30.000	1.00	30.900	3.0	4.18e+005	4.56	Base To Base			
26		262-4	QC 250	Quality Control	******	⊠	250.00	1.00	254.20	1.7	3.21e+006	4.54	Base To Base		······	
27		262-7	QC 750	Quality Control	*******	☒	750.00	1.00	762.50	1.7	8.31e+006	4.50	Base To Base			
28		257-1 264-1	Mobile Phase	Unknown	*******		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			
30		264-2	LT Stb 30 ng/mL LT Stb 30 ng/mL	Unknown			N/A N/A	1.00 1.00	30,959 32,234	N/A	4.19e+005	4.54	Base To Base			
31		264-3	LT Stb 30 ng/mL	Unknown			N/A	1.00	32.234	N/A N/A	4.36e+005 4.40e+005	4.51 4.49	Base To Base	<u> </u>		
32		264-4	LT Stb 750 ng/mL	Unknown				1.00	771.82	N/A	8.39e+006	4.49	Base To Base Base To Base			
33		264-5	LT Stb 750 ng/mL	Unknown	,		N/A	1.00	749.67	N/A	8.21e+006	4.48	Base To Base	$\vdash \dashv$		
34	I6-534006p\I6-1680.wiff	264-6	LT Stb 750 ng/mL	Unknown			N/A	1.00	759.13	N/A	8.28e+006	4.46	Base To Base	ᅡ片ㅣ		
35	16-534006p\l6-1681.wiff	257-1	Mobile Phase	Unknown	,,,,,,,,,		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			
36		261-2	C 10	Standard		⊠	10.000	1.00	10.104	1.0	1.41e+005	4.43	Base To Base			
37		261-5	C 30	Standard		⊠	30.000	1.00	30,771	2.6	4.16e+005	4.44	Base To Base		······	
38		261-8	C 100	Standard		⊠	100.00	1,00	98.832	-1.2	1.30e+006	4.40	Base To Base			
39			C 300	Standard		⊠	300.00	1.00	298.85	-0.38	3.73e+006	4.43	Base To Base			
40		261-14 261-17	C 500 C 750	Standard Standard	******		500.00	1.00 1.00	489.64 754.30	-2.1	5.79e+006	4.39	Base To Base			
42			C 1000	Standard			750.00 1000.0	1.00	754.36 1028.5	0.58 2.8	8.24e+006 1.03e+007	4.42 4.38	Base To Base			
43		257-1	Mobile Phase	Unknown		⊠	N/A	1.00	No Peak	N/A	0.00e+000	0.00	Base To Base No Peak			l
44		262-2	QC 30	Quality Control	********	⊠			32,115		4.34e+005	4.32	Base To Base			İ
45		262-5	QC 250	Quality Control	,,			1.00	259.31		3.27e+006	4.36	Base To Base		,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	İ
46		262-8	QC 750	Quality Control							8.77e+006		Base To Base			Í
47		257-1	Mobile Phase	Unknown					No Peak		0.00e+000	0.00	No Peak			İ
48		265-1	FT Stb 30 ng/mL	Unknown		······································	N/A	1.00	31.881		4,31e+005	4.31	Base To Base			I
49			FT Stb 30 ng/mL	Unknown				1.00	32.061	N/A	4.33e+005	4.30	Base To Base			ł
50			FT Slb 30 ng/mL	Unknown					32.648	N/A	4.41e+005	4.30	Base To Base		**	ł
51 52			FT Stb 30 ng/mL	Unknown			N/A		32.715		4.42e+005		Base To Base			
53			FT Stb 30 ng/mL	Unknown			N/A	1.00	33.364		4.51e+005	4.29	Base To Base			
54	L		FT Stb 30 ng/mL FT Stb 30 ng/mL	Unknown			N/A	1.00	32.896		4.44e+005	4.27	Valley			
55			FT Stb 30 ng/mL	Unknown			N/A N/A	1.00 1.00			4.58e+005	4.27	Base To Base			
56			FT Sib 30 ng/mL	Unknown			N/A	1.00	32.560		4.46e+005 4.40e+005	4.28 4.28	Base To Base	무나		
57	l		FT Stb 750 ng/mL	Unknown			N/A		784.28		4.40e+005 8.49e+006	4.28	Base To Base Base To Base			
58			FT Stb 750 ng/mL	Unknown							8.46e+006		Base To Base			í
L	·				<u>l</u>								230 IA DU2A	<u> </u>		:

Page 2 of 2

Results Path: \\Lcms	sp03\sciexdata\Projects	s\534006\Bio\Resu	lts\16-534006p1.rdt

	File Name	Sample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record	Analyte Concentration (ng/mL)	Dilution Factor	Calculated Concentration (ng/mL)	%RE	Analyte Peak Area (counts)	Analyte Retention Time (min)	Analyte Integration Type	Record Modified	Sample Annotatio
59	I6-534006p\I6-1705.wilf	265-12	FT Slb 750 ng/mL	Unknown			N/A	1.00	770.35	N/A	8.38e+006	4.23	Base To Base	П	Maria Maria - Ing
0	16-534006p\16-1706.wiff	265-13	FT Slb 750 ng/mL	Unknown			N/A	1.00	768.36	N/A	8.36e+006	4.24	Base To Base	Hā	
1	16-534006p\16-1707.wiff	265-14	FT Stb 750 ng/mL	Unknown			N/A	1.00	764.71	N/A	8.33e+006	4.23	Base To Base		
2	16-534006p\16-1708.wiff	265-15	FT Slb 750 ng/mL	Unknown			N/A	1.00	795.48	N/A	8.58e+006	4.22	Base To Base		······································
3	16-534006p\16-1709.wiff	265-16	FT Slb 750 ng/mL	Unknown			N/A	1.00	789.68	N/A	8.54e+006	4.22	Base To Base		
		265-17	FT Slb 750 ng/mL	Unknown			N/A	1.00	791.67	N/A	8.55e+006	4.22	Base To Base		
		265-18	FT Stb 750 ng/mL	Unknown			N/A	1.00	793.75	N/A	8.57e+006	4.24	Base To Base		
6	16-534006p\16-1712.wiff	257-1	Mobile Phase	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak	П	
7	16-534006p\16-1713.wiff	262-3	QC 30	Quality Control		⊠	30.000	1,00	33.998	13.	4.59e+005	4.22	Base To Base	n	
8	16-534006p\16-1714.wiff	262-6	QC 250	Quality Control		×	250.00	1.00	273.11	9.2	3.43e+006	4.19	Base To Base		
9	16-534006p\16-1715.wiff	262-9	QC 750	Quality Control		⊠	750.00	1.00	850.43	13.	9.02e+006	4.17	Base To Base		
0	16-534006p\16-1716,wiff	257-1	Mobile Phase	Unknown	-		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		
1	16-534006p\16-1717.wiff	261-3	C 10	Standard		⊠	10.000	1.00	10.842	8.4	1.51e+005	4.20	Base To Base		······
2	16-534006p\16-1718.wilf	261-6	C 30	Slandard		⊠	30.000	1.00	32.772	9.2	4.43e+005	4.17	Base To Base		
3	16-534006p\16-1719.wiff	261-9	C 100	Standard		⊠	100.00	1.00	105.86	5.9	1.39e+006	4.18	Base To Base		
		261-12	C 300	Standard		×	300.00	1.00	314.78	4.9	3.91e+006	4.19	Base To Base		
	16-534006p\l6-1721.wiff	261-15	C 500	Standard		⊠	500.00	1.00	508.36	1.7	5.98e+006	4.14	Base To Base		
		261-18	C 750	Standard		×	750.00	1.00	774.67	3.3	8.41e+006	4.16	Base To Base		
7	16-534006p\16-1723.wiff	261-21	C 1000	Standard	***************************************	×	1000.0	1.00	1073.8	7.4	1.06e+007	4.15	Base To Base		Peak splitting factor

Printing Date: Saturday, July 21, 2007



Printing Date: Saturday, July 21, 2007 Printing Time: 10:08:35 AM

240 of 281

-144-

Operator: Sheliy Hollar nalyst Version: 1.4.2 -145-241 of 281

Note:

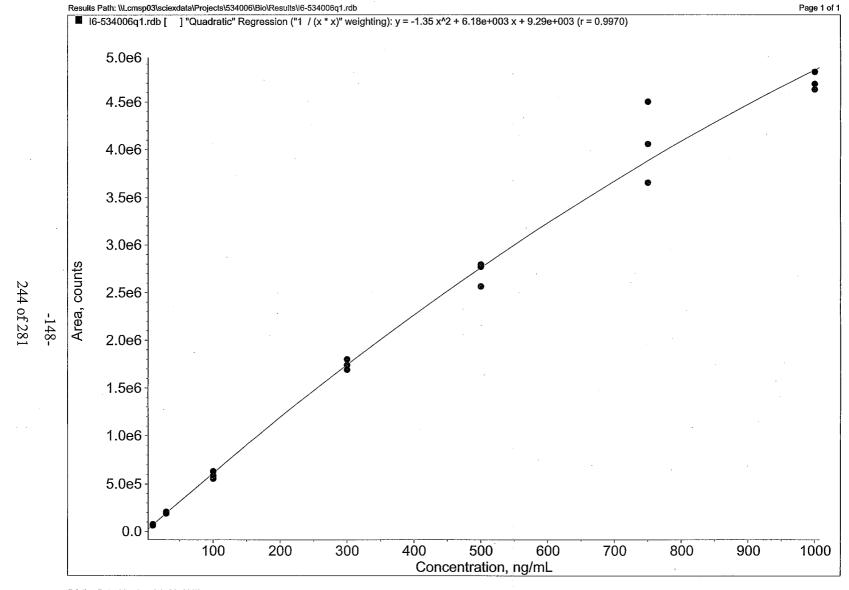
- Urine Long Term Stability
- Urine Freeze Thaw Stability

Study Record Page: 275b to 275c

Res	ults Path: \\Lcmsp0	3\sciex	data\Projects\5340	006\Bio\Resu	Its\I6-53400	)6q1.rc	lb										Page 1 of
	File Name	Sample			Analyte Peak	Use	Analyte	Ditution	Calculated		Analyte Peak	Analyte	Analyte	Record			
13	rile Name	ID	Sample Name	Sample Type	Name	Record	Concentration (ng/mL)	Factor	Concentration (ng/mL)	%RE	Area (counts)	Retention Time (min)	Integration Type	Modified	Sample Annotation	EEA	
1	16-534006q\16-1743.wiff	267-1	Mobile Phase	Unknown		39805200	N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak	37%/5788888 	######################################	Mobile Phase	
2	16-534006q\16-1744.wiff	267-1	Mobile Phase	Unknown	1	<del> </del>	N/A	1.00	No Peak	NA	0.00e+000	0.00	No Peak		·····	Mobile Phase	
3	16-534006q\16-1745.wiff	267-1	Mobile Phase	Unknown	1 -	<b></b>	N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		·····	Mobile Phase	
4	16-534006q\16-1746.wiff	267-1	Mobile Phase	Unknown	1 _		N/A		No Peak	N/A	0.00e+000	0.00	No Peak			Mobile Phase	
5	16-534006q\16-1747.wiff	267-1	Mobile Phase	Unknown	I		N/A		No Peak	N/A	0.00e+000	0.00	No Peak		***************************************	Mobile Phase	
6	16-534006q\16-1748.wilf 16-534006q\16-1749.wilf	267-1 267-1	Mobile Phase Mobile Phase	Unknown		ļ	N/A		No Peak	N/A	0.00e+000	0.00	No Peak			Mobile Phase	
/ B	16-534006q\16-1749.will	267-1	Mobile Phase	Unknown	┦	ļ	N/A		No Peak	N/A	0.00e+000	0.00	No Peak			Mobile Phase	
9	16-534006q\l6-1751.wiff	267-1	Mobile Phase	Unknown	<b>∤</b>	ļ	N/A N/A		No Peak No Peak	N/A N/A	0.00e+000 0.00e+000	0.00 0.00	No Peak No Peak			Mobile Phase	
10	I6-534006q\I6-1752,wiff	267-1	Mobile Phase	Unknown		<del> </del>	N/A		No Peak	N/A	0.00e+000	0.00	No Peak			Mobile Phase Mobile Phase	
11	16-534006q\16-1753.wiff	267-1	Mobile Phase	Unknown	1	<del> </del>	N/A		No Peak	N/A	0.00e+000	0.00	No Peak			Mobile Phase	
12	16-534006q\16-1754.wiff	267-1	Mobile Phase	Unknown	1 -	·····	N/A		No Peak	N/A	0.00e+000	0.00	No Peak			Mobile Phase	
13	16-534006q\16-1755.will	267-1	Mobile Phase	Unknown	† -		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak		······································	Mobile Phase	
14		267-1	Mobile Phase	Unknown	]		N/A		No Peak	N/A	0.00e+000	0.00	No Peak	6		Mobile Phase	
15	16-534006q\16-1757.wiff		Mobile Phase	Unknown	I		N/A		No Peak	N/A	0.00e+000	0.00	No Peak		***************************************	Mobile Phase	
16 17	16-534006q\16-1758.wiff	267-1	Mobile Phase	Unknown	ļ <u> </u>		N/A		No Peak	N∕A	0.00e+000	0.00	No Peak			Mobile Phase	
17 18	16-534006q\16-1759.wiff	267-1 267-1	Mobile Phase	Unknown			N/A		No Peak	N/A	0.00e+000	0.00	No Peak			Mobile Phase	
18 19	16-534006q\16-1760.wiff 16-534006q\16-1761.wiff	267-1 267-1	Mobile Phase Mobile Phase	Unknown Unknown			N/A N/A		No Peak No Peak	N/A N/A	0.00e+000	0,00	No Peak			Mobile Phase	
20	16-534006q\16-1762.wiff	267-1	Mobile Phase	Unknown	<b>∤</b>				No Peak No Peak	1	0.00e+000	0.00	No Peak		***************************************	Mobile Phase	
21	16-534006q\16-1763.wiff	267-1	Mobile Phase	Unknown			N/A N/A		No Peak No Peak	N/A N/A	0.00e+000 0.00e+000	0.00 0.00	No Peak No Peak	무	······································	Mobile Phase	
22	16-534006q\16-1764,wiff	267-1	Mobile Phase	Unknown			N/A		No Peak	N/A	0.00e+000	0.00	No Peak	<u></u>		Mobile Phase Mobile Phase	į
23	16-534006q\16-1765.wiff	267-1	Mobile Phase	Unknown	1 -		N/A		No Peak	N/A	0.00e+000	0.00	No Peak			Mobile Phase	
24	16-534006q\16-1766.wiff	267-1	Mobile Phase	Unknown	<b>!</b>		N/A		No Peak	N/A	0.00e+000		No Peak			Mobile Phase	
25		267-1	Mobile Phase	Unknown	<b>!</b>		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak	Ö		Mobile Phase	
26	16-534006q\16-1768.will	267-1	Mobile Phase	Unknown			N∕A		No Peak	N/A	0.00e+000	0.00	No Peak			Mobile Phase	
27	16-534006q\16-1769.wiff	267-1	Mobile Phase	Unknown					No Peak	N/A	0.00e+000	0.00	No Peak			Mobile Phase	
28		267-1	Mobile Phase	Unknown			N/A		No Peak	NA	0.00e+000	0.00	No Peak			Mobile Phase	
29 30	16-534006q\t6-1771.wiff 16-534006q\t6-1772.wiff	267-1 268-10	Mobile Phase Test Solution	Unknown			N/A		No Peak	N/A	0.00e+000	0.00	No Peak			Mobile Phase	
31	16-534006q\16-1773.wiff	268-10	Test Solution	Unknown			N/A N/A		No Intercept No Intercept	#BAD!	1.04e+007		Base To Base			Test Solution	
32		268-10	Test Solution	Unknown					No Intercept	#BAD! #BAD!	1.03e+007 1.02e+007	4.70 4.71	Base To Base Valley			Test Solution	
33		268-10	Test Solution	Unknown	ļ		N/A		No Intercept	#BAD!			Valley			Test Solution	
34	16-534006q\16-1776.wiff	268-10	Test Solution	Unknown	-				No Intercept	#BAD!			Base To Base			Test Solution Test Solution	
35	16-534006q\16-1777,wiff	268-10	Test Solution	Unknown					No Intercept	#BAD!			Valley			Test Solution	
36	16-534006q\16-1778.wilf	268-10	Test Solution	Unknown			N/A	1.00	No Intercept	#BAD!	1.03e+007		Base To Base		<del></del>	Test Solution	İ
37	16-534006qV6-1779.wilf	268-10	Test Solution	Unknown			N/A	1.00	No intercept	#BAD!	1.04e+007	4.74	Base To Base		***************************************	Test Solution	
38	16-534006q\16-1780.wiff	268-10	Test Solution	Unknown			N/A		No Intercept	#BAD!			Base To Base		·····	Test Solution	
39	16-534006q\16-1781.wiff	268-10	Test Solution	Unknown					No Intercept	#BAD!			Base To Base			Test Solution	İ
40 41	G-534006q\l6-1782.wiff  G-534006q\l6-1783,wiff	267-1 272-1	Mobile Phase Solvent Blank	Unknown Unknown					No Peak	N/A			No Peak			Mobile Phase	
42	16-534006q\16-1784.wilf		Blank Urine	Unknown			N/A N/A		No Peak No Peak	N/A N/A			No Peak			Solvent Blank	
43		272-3	Blank Urine	Unknown			N/A			N/A			No Peak No Peak			Blank Urine Blank Urine	İ
44	16-534006q\16-1786.wiff	272-4	Blank Urine	Unknown					No Peak				No Peak			Blank Urine	
45	16-534006q\16-1787.wiff	267-1	Mobile Phase	Unknown	-				No Peak		0.00e+000		No Peak			Mobile Phase	İ
46	16-534006q\16-1788.wiff	270-1	C 10	Standard			10,000		9.0316				Base To Base			C 10	
47		270-4	C 30	Standard	_		30.000	1.00	29.429	-1.9			Base To Base			C 30	ł
48			C 100	Standard	[ ""	. 🗵			90.510				Base To Base	ᡖ		C 100	
49			C 300	Standard					290.56	-3.1			Base To Base			C 300	
50 51			C 500	Standard					501.88				Base To Base			C 500	
51 52	16-534006q\l6-1793.wiff 16-534006q\l6-1794.wiff		C 750 C 1000	Standard									Base To Base			C 750	
52 53	16-534006q\16-1795.wiff		Mobite Phase	Standard Unknown									Base To Base			C 1000	
54			QC 30	Quality Control					No Peak 32,138				No Peak			Mobile Phase	
55			QC 250	Quality Control						7.1 -13.			Base To Base Base To Base	_무		QC 30	
56			QC 750	Quality Control									Base To Base			QC 250 QC 750	
57			Mobile Phase	Unknown	-					N/A			No Peak			Mobile Phase	
58		273-1	LT SIb 30 ng/mL	Unknown					31.352	N/A			Base To Base			LT SIb 30 ng/mL	
59.		273-2	LT Stb 30 ng/mL	Unknown			WA	1.00	25.687	N/A			Base To Base	-H		LT Stb 30 ng/mL	
60		273-3	LT Stb 30 ng/mL	Unknown								4.72	Base To Base			LT Slb 30 ng/mL	
61			LT Stb 750 ng/mL	Unknown									Base To Base			LT Stb 750 ng/m	
62	16-534006q\16-1804.wiff	2/3-5	LT Stb 750 ng/mL	Unknown		T	WA	1.00	B44.60	N/A	4.26e+006	4.73	Base To Base			LT Slb 750 ng/m	

1978. 1881	File Name	Sample ID	Sample Name	Sample Type	Analyte Peak Name	Use Record		Dilution Factor	Calculated Concentration (ng/mL)	%RE	Analyte Peak Area (counts)	Analyte Retention Time (min)	Analyte Integration Type	Record Modified	Sample Annotation	EEA
			LT Stb 750 ng/mL	Unknown	F		N/A	1.00	881.42	N/A	4.41e+006	4.73	Base To Base	ΪП		LT Stb 750 ng/n
i4	16-534006q\16-1806.wiff		Mobile Phase	Unknown	i	1	N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak	1-1		Mobile Phase
55	16-534006q\16-1807.wiff		C 10	Standard	ī	⊠	10.000	1.00	11.033	10.	7.73e+004	4.72	Base To Base	- H		C 10
6	(6-534006q\16-1808,wiff		C 30	Standard	ī	⊠	30.000	1.00	32.296	7.7	2.07e+005	4.76	Base To Base		<u> </u>	C 30
7	16-534006q\16-1809,will		C 100	Standard	ì	×	100.00	1.00	95.955	4.0	5.89e+005	4.74	Base To Base	T 7	†	C 100
8	16-534006q\16-181D,will		C 300	Standard	i ~~	$\boxtimes$	300.00	1.00	299.35	-0.22	1.74e+006	4.73	Base To Base	1 5		C 300
59	16-534006q\t6-1811.wiff		C 500	Standard	1	⊠	500.00	1.00	507.04	1.4	2.79e+006	4.76	Base To Base		<u> </u>	C 500
'O :	<del>1</del>		C 750	Standard	i ·····	⊠	750.00	1.00	907.23	21.	4.50e+006	4.76	Base To Base	1 5	<del> </del>	C 750
		1	C 1000	Standard	1	⊠	1000.0	1.00	958.52	-4.1	4.69e+006	4.75	Base To Base	1 -	·	C 1000
	1		Mobile Phase	Unknown	i · · · · · · · · · · · · · · · · · · ·		N/A	1,00	No Peak	N/A	0.00e+000	0.00	No Peak	1 5		Mobile Phase
			QC 30	Quality Control	i	⊠	30.000	1.00	31.410	4.7	2.02e+005	4.75	Base To Base	15	†	QC 30
	I	1	QC 250	Quality Control	1	⊠	250.00	1.00	232.45	-7.0	1.37e+006	4.74	Base To Base		<u> </u>	QC 250
			QC 750	Quality Control		☒	750.00	1.00	711.35	-5.2	3.72e+006	4.73	Base To Base			QC 750
			Mobile Phase	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			Mobile Phase
7		274-1	FT Stb 30 ng/mL	Unknown			N/A	1.00	32.932	N/A	2.11e+005	4.74	Base To Base	1 5	l	FT Stb 30 ng/m
			FT Stb 30 ng/mL	Unknown	<b></b>		N/A	1.00	29.877	NA	1.93e+005	4.77	Base To Base	T 17		FT Stb 30 ng/m
9			FT Stb 30 ng/mL	Unknown			N/A	1.00	29.460	N/A	1.90e+005	4.73	Base To Base			FT Stb 30 ng/m
0			FT Slb 750 ng/mL	Unknown			N/A	1.00	830.31	N/A	4.21e+006	4.72	Base To Base			FT Stb 750 ng/r
			FT Stb 750 ng/mL	Unknown	•		N∕A	1.00	769.97	N/A	3.97e+006	4.73	Valley		······································	FT Stb 750 ng/r
			FT Stb 750 ng/mL	Unknown			N/A	1.00	876.16	N/A	4.39e+006	4.72	Base To Base	1 H	1	FT Stb 750 ng/r
			Mobile Phase	Unknown			N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak			Mobile Phase
			QC 30	Quality Control	••••	⊠	30.000	1.00	28.890	-3.7	1.87e+005	4.75	Base To Base	1 6		QC 30
	16-534006q\t6-1827.wiff		QC 250	Quality Control		⊠	250.00	1.00	233.37	-6.7	1.38e+006	4.75	Base To Base			QC 250
rivineisi.	***************************************		QC 750	Quality Control		⊠	750.00	1.00	749.42	-0.077	3.88e+006	4.75	Valley		·	QC 750
			Mobile Phase	Unknown	*****		N/A	1.00	No Peak	N/A	0.00e+000	0.00	No Peak	T	ļ	Mobile Phase
			C 10	Standard		×	10.000	1.00	9.6527	-3.5	6.88e+004	4.74	Base To Base			C 10
			C 30	Standard		×	30.000	1.00	31.740	5.8		4.75	Base To Base	<del>                                     </del>	<del> </del>	C 30
		270-9	C 100	Standard		×	100.00	1.00	103.21	3.2	6.32e+005	4.74	Base To Base	1-5-	İ	C 100
		270-12	C 300	Standard	*****	×	300.00	1.00	310.58	3.5			Base To Base		<del></del>	C 300
		270-15	C 500	Standard		☒	500.00	1.00	459,59	-8.1			Base To Base			C 500
		270-18		Standard		×	750.00	1.00	696.11	-7.2	3.66e+006		Base To Base	<u> </u>		C 750
4	16-534006q\16-1836.wiff	270-21	C 1000	Standard			1000.0	1.00	994.27	-0.57			Base To Base	H		C 1000





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Operator: Shelly Hollar

## APPENDIX F

Toxicokinetic Report [

## STUDY TITLE

## PHARMACOKINETIC (IN BLOOD) AND EXCRETION STUDY OF [ ] IN RATS

## **REPORT TITLE**

# PHARMACOKINETICS OF [ ] IN SERUM AND URINE FOLLOWING A SINGLE INTRAVENOUS DOSE TO RATS

## **REPORT DATE**

26 September 2007

FINAL REPORT

<u>Sponsor</u>	<b>Testing Facility</b>
[ ]	[ ]

## TABLE OF CONTENTS

		Page	
LIST	OF TA	ABLES3	
LIST	OF FIG	GURES3	
1.0	SUM	MARY4	
2.0	INTE	RODUCTION6	
3.0	EXP	ERIMENTAL7	
	3.1	Data Processing7	
	3.2	Bioanalysis7	
	3.3	Toxicokinetic Evaluation & Statistical Analysis	÷
4.0	RESU	ULTS AND DISCUSSION9	
	4.1	Serum Concentration Data9	
	4.2	[ ] Elimination in Urine	
	4.3	[ ]EEA Pharmacokinetics	15
5.0	CON	CLUSIONS17	
6.0	RESF	PONSIBLE PERSONNEL	

## LIST OF TABLES

<u>Table</u>	<u>Title</u>	<u>Page</u>
Table 1.	Mean ± SD Concentrations of [ ] in Serum of Male Rats following Intravenous Administration of 10 mg [ ]/kg	10
Table 2.	Mean ± SD Amounts of [ ] in Urine following Intravenous Administration of 10 mg [ ]/kg to Male and Female Rats	12
Table 3.	Pharmacokinetic Parameters for [ ]in Serum following Intravenous Administration of 10 mg [ ]/kg to Male and Female Rats	15
Table 4.	Pharmacokinetic Parameters for [ ] in Urine following Intravenous Administration of 10 mg [ ]/kg to Male and Female Rats	16
	LIST OF FIGURES	
<u>Figure</u>	<u>Title</u>	<u>Page</u>
Figure 1.	Mean + SD Concentrations of [ ] in Serum following Intravenous Administration of 10 mg [ ]/kg to Male and Female Rats	11
Figure 2.	Cumulative Percent of Dose Eliminated in Urine following Intravenous Administration of 10 mg [ ]/kg to Male and Female Rats	13
Figure 3.	Mean Amount of [ ] Dose Remaining to Be Eliminated in Urine of Male and Female Rats following Intravenous Administration of 10 mg	1.4

#### 1.0 SUMMARY

One group of nine male and nine female Crl:CD®(SD) rats received a single intravenous (bolus) injection of difluoro [

] at a dosage of 10 mg/kg at a dosage volume of 5 mL/kg. Blood samples were collected from three animals/sex/group at 0 (prior to dosing), 2, 10, 20, and 30 minutes and 1, 3, 5, 7, 24, and 48 hours after dosing.

For the urine excretion phase, one group of three male and three female Crl:CD[®](SD) rats received a single intravenous (bolus) injection of [ ] at a dosage level of 10 mg/kg at a dosage volume of 5 mL/kg. Urine was collected from each animal over the following intervals: 0-6, 6-12, and 12-24 hours post-dosing. After each urine collection, cages were rinsed and the rinses were collected separately for analysis.

The concentration of [ ] in the serum, urine, and cage rinse samples was measured using a validated LC-MS/MS method. The serum concentration immediately following the intravenous dose was estimated based on a regression analysis of the measured values. The mean concentrations in serum and mean amounts excreted in urine plus cage rinse were used for pharmacokinetic analysis.

The pharmacokinetic parameters for [ ] are summarized in the following table:

PHARMAC	OKINETI	C RESULTS I	OR[]						
[]			URINE†						
10 mg/kg Intravenous Dose	C ₀ * (ng/mL )	AUC _{0-∞} (ng×h/mL)	K _{e1} (h ⁻¹ )	Half- life** (h)	Cl (L/h×kg)	V _d (L/kg)	K _{el} (h ⁻¹ )	Half- life** * (h)	% of Dose Elimi- nated
Males Females	69775 102835	373393 53137	0.127 0.074	5.4 9.4	0.0268 0.188	0.210 2.55	0.215 0.392	3.2 1.8	67.3 64.0

^{*} Values were estimated.

After a single intravenous dose of [ ] at 10 mg/kg, systemic exposure (AUC $_{0-\infty}$ ) to [ ] for male rats was almost 7-fold higher than for female rats. [ ] appeared to remain mostly in the circulation in male rats (apparent volume of distribution about 0.2 L/kg), but to have extensive tissue distribution in female rats (apparent volume of distribution of more than 2.5 L/kg). The terminal elimination phase for [ ] in serum had a half-life of 9.4 and 5.4 hours for female and male rats, respectively. The half-life for [ ] in urine was 1.8 and 3.2 hours, for female and male rats respectively. Nevertheless, the percent of [ ] dose eliminated over 24 hours post-dosing in the urine of male rats and female rats was similar (approximately 65%). This can be explained by the lower amounts of [ ] available for urinary clearance in the circulation of female rats compared to male rats as suggested by the differences in apparent volume distribution.

^{**}For the terminal elimination phase.

^{***}For urinary elimination.

[†]Urine plus cage rinse

### 2.0 INTRODUCTION

One group of nine male and nine female Crl:CD®(SD) rats received a single intravenous (bolus) injection of difluoro [

] at a dosage of 10 mg/kg at a dosage volume of 5 mL/kg. Blood samples were collected from three animals/sex/group at 0 (prior to dosing), 2, 10, 20, and 30 minutes and 1, 3, 5, 7, 24, and 48 hours after dosing. Blood samples (approximately 0.5 mL) were collected via a retro-orbital sinus into tubes containing no anticoagulant while the animal was under isoflurane anesthesia. Serum was separated using a refrigerated centrifuge. Samples were stored at approximately -20°C until transferred to the Analytical Chemistry Department [ ], then stored at -70°C until analysis.

For the urine excretion phase, one group of three male and three female Crl:CD®(SD) rats received a single intravenous (bolus) injection of [ ] at a dosage level of 10 mg/kg at a dosage volume of 5 mL/kg. Urine was collected from each animal over the following intervals: 0-6, 6-12, and 12-24 hours post-dosing. After each urine collection, cages were rinsed and the rinses were collected separately for analysis. Urine samples were maintained on wet ice during collection. Samples were stored at approximately -70°C until transferred to the Analytical Chemistry Department of [ ], then stored at -20°C until analysis..

The concentration of [ ] in serum, urine and cage rinse samples were measured by the Analytical Chemistry Department at [ ] using a validated

LC-MS/MS method. The serum concentration immediately following the intravenous dose was estimated based on a regression analysis of the measured values. The mean concentrations in serum and mean amounts excreted in urine (plus cage rinse) were used for pharmacokinetic analysis.

#### 3.0 EXPERIMENTAL

## 3.1 Data Processing

All calculations were performed using Microsoft[®] Excel 2002 on a Microsoft[®] Windows XP Professional platform. Graphical presentations were created using SigmaPlot 2004 for Windows Version 9.0.

## 3.2 Bioanalysis

Concentration of [ ] in serum, urine, and cage rinse samples were measured using a validated LC-MS/MS method by the Analytical Chemistry Department at [

]. A detailed description of the analytical method and the results for each sample may be found in Appendix E. The lower limit of quantitation (LLOQ) was 10 ng/mL for serum, urine and cage rinse.

## 3.3 Toxicokinetic Evaluation & Statistical Analysis

In the calculation of the toxicokinetic parameters, samples were assigned a value of zero if the concentration was below the LLOQ. All toxicokinetic parameters were calculated

# 534006

from the mean serum or urine (plus cage rinse) concentration data as indicated in the following table:

C ₀	The estimated concentration of the compound in serum immediately following intravenous administration. The values were set equal to the y-intercept of the linear regression based on the log concentration of the mean values from 2 to 20 minutes post-dosing.
AUC _{0-t}	The area under the serum concentration vs. time curve from the time of dosing to $C_{last}$ (the last mean serum concentration $> 0$ ). The values were calculated by linear trapezoidal summation using the equation:
	$AUC_{0-t} = \Sigma (0.5 \cdot (y_1 + y_2) \cdot \Delta t)$
	where $y_1$ and $y_2$ are successive serum concentrations and $\Delta t$ is the sampling interval, in hours, between $y_1$ and $y_2$ .
AUC _{0-∞}	The estimate of the area under the serum concentration vs. time curve from time of dosing to infinity. The values were calculated using the formula: $AUC_{0-\infty} = AUC_{0-t} + (C_{last}/K_{el})$ where AUCa, and $C_{t-t}$ were defined previously, and $K_{t-t}$ is
	where $AUC_{0-t}$ and $C_{last}$ were defined previously, and $K_{el}$ is defined subsequently.
K _{el}	The terminal elimination rate constant for the compound in serum or urine. The values were calculated using the equation: $K_{el} = -\ln[10] \ x \ b$ where b is the slope of the least-squares linear regression line of
	the log serum concentrations from 7 to 48 hours post-dosing or the log ARE from 0 to 12 hours post-dosing and ARE is defined subsequently.
Half-life	The half-life for the compound in serum or the half-life of urinary elimination. The values were calculated using the formula:
	$Half-life = -ln[0.5]/K_{el}$
	where K _{el} is defined previously.

Cl	The apparent systemic clearance for the compound in serum. The values were calculated using the formula: $Cl = Dosage/AUC_{0-\infty}$ where $AUC_{0-\infty}$ is defined previously.
$ m V_d$	The apparent volume of distribution for the compound in serum. The values were calculated using the formula: $V_d = \text{Cl/K}_{el}$ where Cl and Kel are defined previously.
ARE	The amount remaining to be eliminated in urine. The values were calculated using the formula:  ARE = Total amount eliminated – Amount eliminated in previous interval(s)
Urinary Elimination as % Dose	The total amount eliminated in urine expressed as a percentage of the analyte dose. The value was calculated using the equation:  Total as % Dose = 100 × ARE at 0 h/(Mean BW × Dosage)  where ARE is defined previously and BW is the mean body weight used to calculate the administered dose. Mean body weights were obtained from Table 6 of the main report.

# 4.0 RESULTS AND DISCUSSION

# 4.1 Serum Concentration Data

The concentration of [ ] in individual serum samples can be found in the tables of the bioanalytical report (Appendix E).

Mean serum concentrations of [ ] following a single intravenous dose at 10 mg [ ]/kg to male and female rats are presented in Table 1 and illustrated in Figure 1. The dose for Animal No. 46669 (male) appeared to be delivered extravascularly; mean and

# 534006

SD serum [ ] concentrations for the males are presented with and without the data for this animal.

Table 1. Mean ± SD Concentrations of [ ] in Serum of Male Rats following Intravenous Administration of 10 mg [ ]/kg

Hours	Males		Mal	Males*		Females	
Post- Dosing	Mean (ng/mL)	SD	Mean (ng/mL)	SD	Mean (ng/mL)	SD	
Pre-dose	0.00	0.00	0.00	0.00	0.00	0.00	
$C_0^{**}$	69775	N/A	93101	N/A	102835	N/A	
0.0333	67655	44748	93115	10756	94742	5213	
0.167	69129	8686	69129	10756	50222	5972	
0.333	62353	5666	62353	5666	30007	2019	
0.5	43079	20222	54631	4142	237-15	3816	
1	63725	17015	63725	17015	14424	3930	
3	24953	11314	24953	11314	882	223	
5	15750	6318	13287	6590	361	248	
7	17372	9052	17372	9052	220	170	
24	192	177	192	177	11.8	1.71	
48	73.1	90.9	20.6	3.33	8.98	7.78	

^{*}Data for Animal No. 46669 at 0.033, 0.5, 5 and 48 hours post-dosing were excluded from the calculation of mean and SD values.

^{**}Estimated value. N/A = not applicable

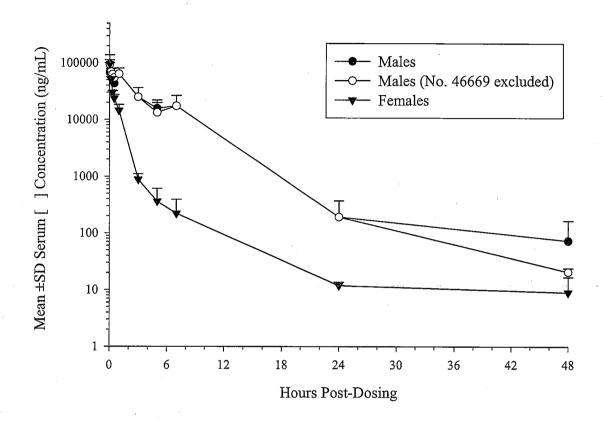


Figure 1. Mean + SD Concentrations of [ ] in Serum following Intravenous Administration of 10 mg [ ]/kg to Male and Female Rats

Concentration of [ ] in serum of male and female rats after single intravenous administration of [ ] at 10 mg/kg was measurable up to 48 hours post-dosing. Elimination of [ ] in serum appeared to multi-phasic in both male and female rats; however, the temporal trends in serum concentration profiles differed between the genders. In male rats, serum [ ] concentration remained relatively constant through 1 hour post-dosing, decreased gradually from 1 to 7 hours post-dosing and then decreased

further from 7 through 48 hours post-dosing. In female rats, serum [ ] concentration decreased rapidly through 3 hours post-dosing, decreased more gradually from 3 to 24 hours post-dosing and then decreased only slightly from 24 through 48 hours post-dosing.

# 4.2 [ ] Elimination in Urine

The concentration of [ ] in individual urine samples can be found in the tables of the bioanalytical report (Appendix E).

Mean amounts of [ ] eliminated in urine following a single intravenous dose at 10 mg [ ]/kg to male and female rats are presented in Table 2. The cumulative percent of [ ] dose eliminated in urine is illustrated in Figure 2 and the amounts of [ ] remaining to be eliminated in urine are illustrated in Figure 3.

Table 2. Mean  $\pm$  SD Amounts of [ ] in Urine following Intravenous Administration of 10 mg [ ]kg to Male and Female Rats

	Male	S	Females	
Hours Post-Dosing	Mean (µg)	SD	Mean (μg)	SD
0-6	878	227	773	132
6-12	571	89.8	300	216
12-24	119	22.2	9.80	3.04

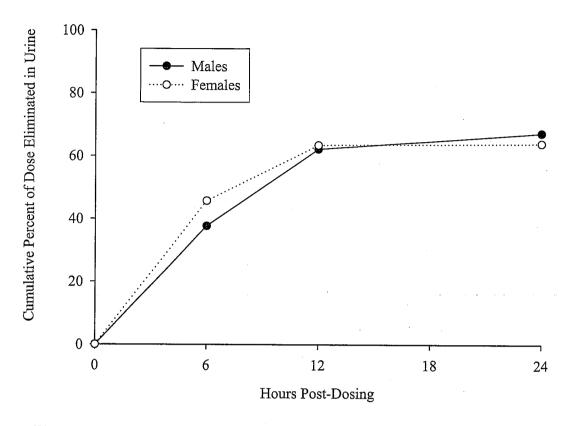


Figure 2. Cumulative Percent of Dose Eliminated in Urine following Intravenous Administration of 10 mg [ ]/kg to Male and Female Rats

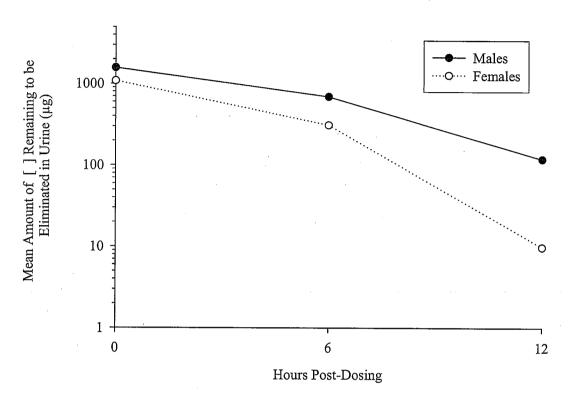


Figure 3. Mean Amount of [ ] Dose Remaining to Be Eliminated in Urine of Male and Female Rats following Intravenous Administration of 10 mg [ ]/kg

The amount of [ ] recovered in the urine accounted for approximately 67% of the administered dose for male rats and about 64% for female rats. Most of the [ ] dose was eliminated in urine over 12 hours post-dosing in both genders. The elimination of [ ] in the urine of male rats appeared to be mono-exponential; in female rats, elimination of [ ] in urine appeared not to be log-linear.

# 4.3 [ ] Pharmacokinetics

The pharmacokinetic parameters for [ ] in serum of male and female rats are presented in Table 3.

Table 3. Pharmacokinetic Parameters for [ ] in Serum following Intravenous Administration of 10 mg [ ]/kg to Male and Female Rats

	AUC _{0-t} (ng×h/mL)	AUC _{0-∞} (ng×h/mL)	K _{el} (h ⁻¹ )*	Half-life (h)*	Cl (L/h×kg)	V _d (L/kg)
Males	372819	373393	0.127	5.4	0.0268	0.210
(Males)**	(373629)	(373758)	(0.160)	(4.3)	(0.0268)	(0.168)
Females	53015	53137	0.0737	9.4	0.188	2.55

^{*} For the terminal elimination phase.

The pharmacokinetic parameters for [ ] differed between the genders. Systemic exposure ( $AUC_{0-\infty}$ ) for male rats was almost 7-fold higher than for female rats. This may be attributable to gender differences in the tissue distribution of [ ] in rats. In male rats, [ ] appeared to remain mostly in circulation (apparent volume of distribution about 0.2 L/kg). In female rats, [ ] appeared to have extensive tissue distribution (apparent volume of distribution of more than 2.5 L/kg). The terminal elimination phase for [ ] in serum had a half-life of 9.4 and 5.4 hours, for female and male rats respectively. Exclusion of the serum concentration data for Animal No. 46669 had negligible effect on most of the pharmacokinetic parameters for male rats.

^{**} Data for Animal No. 46669 were excluded from the calculation of the pharmacokinetic parameters.

Pharmacokinetic parameters for [ ] in urine following a single intravenous dose at 10 mg [ ]/kg to male and female rats are presented in Table 4.

Table 4. Pharmacokinetic Parameters for [ ] in Urine following Intravenous Administration of 10 mg [ ]/kg to Male and Female Rats

	Total Eliminated as a % of Dose	Elimination Rate Constant (1/h)*	Half-life (h)*
Males Females	67.3 64.0	0.215 0.392	3.2 1.8

^{*}Based on 0-12 h.

The half-life for [ ] in urine was 1.8 and 3.2 hours, for female and male rats respectively. Nevertheless, the percent of [ ] dose eliminated over 24 hours post-dosing in the urine of male rats and female rats was similar (approximately 65%). This can be explained by the lower amounts of [ ] available for urinary clearance in the circulation of female rats compared to male rats as suggested by the differences in apparent volume distribution.

#### 5.0 CONCLUSIONS

After a single intravenous dose of [ ] at 10 mg/kg, systemic exposure (AUC $_{0-\infty}$ ) to [ ] for male rats was almost 7-fold higher than for female rats. [ ] appeared to remain mostly in the circulation in male rats (apparent volume of distribution about 0.2 L/kg), but to have extensive tissue distribution in female rats (apparent volume of distribution of more than 2.5 L/kg). The terminal elimination phase for [ ] in serum had a half-life of 9.4 and 5.4 hours for female and male rats, respectively. The half-life for [ ] in urine was 1.8 and 3.2 hours, for female and male rats respectively. Nevertheless, the percent of [ ] dose eliminated over 24 hours post-dosing in the urine of male rats and female rats was similar (approximately 65%). This can be explained by the lower amounts of [ ] available for urinary clearance in the circulation of female rats compared to male rats as suggested by the differences in apparent volume distribution.

		ä	

# APPENDIX G

Study Protocol

#### PROTOCOL AMENDMENT

# A. Title of Study:

Pharmacokinetic (in Blood) and Excretion Study of

#### B. Protocol Modification:

1) 3 STUDY SCHEDULE:

Proposed Experimental Termination (Completion) Date:

April 5, 2007 (last bioanalytical analysis)

#### 2) 4.1 Identification:

#### 8.5 Toxicokinetics for Elimination:

- 3) In accordance with the study director notification dated December 11, 2006, serum and urine samples for toxicokinetic evaluation will be stored at approximately -20°C or lower.
- 5) Also in accordance with the study director notification dated May 3, 2007, the last sentence of this section of the protocol is amended to read as follows:

Subsequently, pertinent toxicokinetic parameters, such as  $C_{max}$ , AUC, and elimination half-life, will be determined as data permit, following a single dose of the test material.

#### 6) 8.6 Anatomic Pathology:

In accordance with the study director notification dated December 13, 2006, this section is added to the protocol. A gross macroscopic examination will be performed on any animals found dead or euthanized *in extremis*. Animals euthanized *in extremis* will be euthanized by  $\mathrm{CO}_2$  inhalation. No tissues will be collected.

# C. Reason for Protocol Modification:

- 1) Addition of Experimental Termination (Completion) date.
- 2) Correction of typographical error.
- 3) Clarification of storage conditions for serum and urine samples.
- 4) Addition of method for calculation of total
- 5) Clarification of wording for a single test material.
- 6) Addition of gross necropsy for animals found dead or euthanized in extremis.

28 June, 2007 Date

22 June 2007

22 Jone 200>



# PROTOCOL

# PHARMACOKINETIC (IN BLOOD) AND EXCRETION STUDY OF

Submitted To:

#### 1 OBJECTIVE:

The objective of this study is to evaluate the pharmacokinetic (in blood) and excretion profiles of the test article in rats.

This protocol has been designed and the study will be conducted in compliance with the U.S. Environmental Protection Agency, 40 CFR Part 792, and the Organisation for Economic Cooperation and Development [C(97)186/Final] Good Laboratory Practice (GLP) Regulations. This study will be performed according to the protocol dard Operating Procedures.

#### 2 PERSONNEL INVOLVED IN THE STUDY:

#### 2.1 Sponsor Representative:

2.2

2,3

2.4

#### 4 TEST ARTICLE DATA:

#### 4.1 Identification:

#### 4.2 Lot Number:

To be provided by the Sponsor.

#### 4.3 Purity:

99.0%. The purity will be considered to be 100% for the purpose of dosage calculations.

#### 4.4 Stability:

The test article is considered to be stable under the storage conditions provided by the Sponsor.

#### 4.5 Physical Description:

To be document

#### 4.6 Storage Conditions:

To be provided by the Sponsor.

#### 4.7 Reserve Samples:

Retention samples will be collected and stored in accordance with Standard Operating Procedures.

#### 4.8 Personnel Safety Information:

To be provided by the Sponsor. It is the responsibility of the Sponsor to notify the testing facility of any special handling requirements for the test article. A material safety data sheet (MSDS) should accompany the test article upon arrival at the laboratory.

#### 4.9 Vehicle:

Sterile water for injection.

#### 5 TEST SYSTEM:

#### 5.1 Species:

Rat

#### 5.2 Strain:

Sprague-Dawley Crl:CD®(SD)

#### 5.3 Source:

Charles River Laboratories, Inc. (Facility to be documented in the study records)

#### 5.4 Number on Study:

Fifteen (15) animals of each sex will be ordered. Twelve males and 12 females will be placed on study. Animals not utilized on study will be deemed as part of the stock colony or euthanized by  $CO_2$  inhalation and discarded.

#### 5.5 Approximate Age and Weight:

Animals will be approximately 5 to 7 weeks of age when received and 7 to 9 weeks of age at initiation of dosing. Body weight will be approximately 200 to 350 grams at initiation of dosing.

#### 5.6 Identification System:

The animals will be uniquely identified by a metal eartag displaying the animal number. Individual cage cards will be affixed to each cage and will display the animal number, group number, study number, dosage level and sex of the animal.

#### 5.7 Justification for Selection and Number on Study:

This species and strain of animal is recognized to be appropriate for subchronic toxicity studies. The Sprague-Dawley rat will be used because it is a widely used strain for which significant historical control data are available. This number of animals is considered to be the minimum required for meaningful interpretation of the data and fulfillment of agency requirements.

#### 6 SPECIFIC MAINTENANCE SCHEDULE:

#### 6.1 Animal Housing:

Animals will be housed in a study-dedicated, environmentally controlled room three per cage by sex in clean, suspended, wire-mesh cages for approximately two to four days following receipt. If the number of animals received does not allow for all animals to be housed three per cage by sex, then some animals will be pair-housed by sex. Thereafter, all animals will be housed individually. The cages will be clevated above cage-board or other suitable material which will be changed in the times each week. The facilities at

fully accredited by the Association for

Laboratory Animal Care International (AAALAC International).

#### 6.2 Environmental Conditions:

Controls will be set to maintain an average daily temperature of  $71 \pm 5^{\circ}$ F ( $22 \pm 3^{\circ}$ C) and an average daily relative humidity of  $50 \pm 20\%$ . Temperature and relative humidity will be monitored continuously. Data for these two parameters will be scheduled for automatic collection on an hourly basis. Fluorescent lighting controlled by light timers will provide illumination for a 12 hour light/dark photoperiod. Temporary adjustments to the light/dark cycles may be made to accommodate protocol specified activities. The ventilation rate will be set at a minimum of 10 room air changes per hour, 100% fresh air.

#### 6.3 Drinking Water:

Reverse osmosis-purified water will be available ad libitum. Filters servicino the automatic watering system are changed regularly according to Standard Operating Procedures. The municipal water supplying the labor is analyzed according to utine basis to assure that contaminants are not present in concentrations that would be expected to affect the outcome of the study.

#### 6.4 Basal Diet:

PMI Nutrition International, LLC Certified Rodent LabDiet[®] 5002 (meal) will be offered ad libitum during the study. Periodic analyses of the certified feed are performed by the manufacturer to ensure that heavy metals and pesticides are not present at concentrations that would be expected to affect the outcome of the study. Results of the analyses are provided to

change

#### 7 EXPERIMENTAL DESIGN:

#### 7.1 Animal Receipt and Quarantine:

Each animal will be inspected by a qualified technician upon receipt. Animals judged to be in good health will be placed immediately in acclimation for at least seven days. All animals will be weighed and assigned a permanent animal number. During the acclimation period, each animal will be observed twice daily for changes in general appearance and behavior. There will be a pretreatment week (as part of the acclimation period) during which body weights and food consumption will be recorded and general health will be monitored, but the rats will not be dosed. All animals will receive a detailed physical examination at the initiation of pretest and at the time of animal selection for randomization.

#### 7.2 Randomization:

At the conclusion of the acclimation period, animals judged to be suitable for testing will be assigned to groups at random, based on body weight stratification into a block design, using a computer program. A printout containing the animal numbers and individual group assignments will be generated. Animals will then be arranged into the groups according to the printout. Body weights at randomization will be within ± 20% of the mean for each sex.

#### 7.3 Route and Rationale of Test Article Administration:

The route of administration will be intravenous since this is an acceptable route of administration to assess systemic exposure.

#### 7.4 Organization of Test Groups. Dosage Levels and Treatment Regimen:

#### 7.4.1 Organization of Test Groups:

The dosage levels will be determined from the results of previous studies and will be provided by the Sponsor Representative after consultation with the The following diagram presents the study group arrangement.

Pharmacokinetic (Blood Collection) Groups:

Group Number	Treatment	Dosage Dosage Level Concentration		Dosage Volume	Number of Animals	
		(mg/kg)	(mg/mL)	(mL/kg)	Males	Females
1		10	2	5	9	9

#### Excretion (Urine Collection) Groups:

Group Number	Treatment	Dosage Level	Dosage Concentration	Dosage Volume	Number of Animals	
Number		(mg/kg)	(mg/mL)_	(mL/kg)	Males	Females
1		, 10	2	5	3	3

Data for pharmacokinetic groups and excretion groups will be collected in separate computer protocols.

#### 7.4.2 Vehicle:

Sterile water for injection will be used as the vehicle.

# 7.4.3 Treatment Regimen:

Animals will be appropriately restrained and administered dosing solutions by a slow bolus intravenous injection (sterile needle and syringe) via a lateral tail vein. A constant volume of 5 mL/kg will-be used. The treatment period will be 1 day. Day 0 will be the day of dosing.

#### 7.5 Preparation and Analysis of Test Article Preparations:

#### 7.5.1 Test Article Preparation:

The test article will be prepared for dosing as weight-to-volume mixtures in a vehicle and filter-sterilized in a laminar flow hood. No correction for purity will be made. A complete description of the method of test article preparation will be documented in the Study records and described in the final report. Test article formulations will be prepared within 1 week of use for dosing and stored refrigerated. The formulation will be removed from storage and allowed to remain at room temperature for at least 1 hour before dosing.

# 7.5.2 Homogeneity and Stability of Test Article Formulations:

Homogeneity assessments will not be performed, as the formulations are solutions. Analyses to demonstrate the stability of the test article formulation for the expected period of refrigerated storage between formulation and dosing will be conducted before the initiation of dosing.

#### 7.5.3 Concentration Analysis:

Concentration will be confirmed during the dosing period. Samples will be drawn from the test article formulation. These will be submitted to

the

nalyzed for

test article concentration using a validated method.

The Analytical Chemistry report will be appended to the final report for this study.

#### 8 EXPERIMENTAL OBSERVATIONS:

#### 8.1 Viability Observations:

All animals will be observed for mortality/moribundity twice daily, once in the morning and once in the afternoon. Moribund animals will be euthanized to ensure that tissues will not be lost due to autolysis.

#### 8.2 Detailed Physical Examination:

All animals will receive a detailed physical examination at least once during the pre-treatment period. Animals without signs will be noted individually.

# 8.3 Individual Body Weights:

Individual body weights will be recorded during acclimation, at pretest initiation, at randomization and on Day 0.

#### 8.4 Individual Food Consumption:

Individual food consumption will be recorded during the pretreatment period only.

#### 8.5 Toxicokinetics for Elimination:

Blood samples will be obtained from the blood collection groups for determination of concentration of the erum at the time points outlined in the following table:

Intervals .	1 Day 0
Time points post- dosing	<ul> <li>Prior to a dosing and approximately 2, 10, 20 and 30 minutes and 1, 3, 5, 7, 24 and 48 hours after dosing</li> <li>Clock times of collection recorded</li> </ul>
Number of Animals	<ul> <li>3 animals/sex bled per time point</li> <li>Each animal sampled no more than three times in a 24 hour period (unless a terminal collection followed by euthanasia).</li> </ul>
Sample Collection	Retro-orbital sinus under isoflurane (inhalation) anesthesia.
Target Blood Volumes	0.5 mL/time point     collect into non-chilled sampling tubes
Anticoagulant	None.
Sample Handling	Samples will be allowed to clot at room temperature, after which they will be kept chilled (ice water bath, as appropriate) after collection and during processing.
Serum Preparation	Beckman 6R centrifuge 2400-2700 rpm at -4°C
Aliquots	Recover all serum possible and place in Nunc [©] plastic vials.
Label information	Study number, dose group, animal number, sample type, date of collection, time of collection.
Storage	Approximately -20°C until analysis     Time placed in freezer recorded

Moribund animals will be euthanized by CO₂ inhalation. Animals found dead or euthanized in extremis after start of dosing will be examined to determine possible cause of death and discarded. Following the final blood collection, all animals will be euthanized by carbon dioxide inhalation and discarded.

Urine collection animals will be transferred into plastic metabolism cages for urine collection following dosing. Urine will be collected on wet ice over the following intervals: 0-6, 6-12 and 12-24 hours post-dosing. The volume of each urine sample will be recorded, after which the urine samples will be frozen with minimal delay in a freezer set to maintain temperature of approximately -20°C until preparation for analysis. Following the final urine collection, all animals used for urine collection will be euthanized by carbon dioxide inhalation and discarded.

During method validation, stability of assessed in processed samples, during long-term frozen storage (-20°C), after short-term (at least 4-

hour) room temperature storage and during the freeze-thaw process. At least triplicate samples will be analyzed and the mean response will be compared against that of the freshly-prepared samples (the sample response may be analyzed as concentration, peak area, or peak area ratio, etc, depending on the stability analysis). If a significant degradation (>15% reduction in the mean response) occurs under any of the tested conditions, then special precautions will be taken.

Serum and urine samples will be analyzed for EEA concentration, by the Analytical Chemistry Department at a validated LC/MS/MS method.

Subsequently, pertinent toxicokinetic parameters, such as  $C_{max}$ , AUC, and elimination half-life, will be determined as data permit, for each of the test articles following a single dose of the test material.

#### 9 STATISTICAL METHODS:

No statistical test will be performed.

#### 10 QUALITY ASSURANCE:

The study will be audited by the assure compliance with the study protocol and protocol amendments, WIL Standard Operating Procedures and the appropriate provisions of the U.S. EPA TSCA and FIFRA Good Laboratory Practice Standards published in the Federal Register (40 CFR Part 792 and 40 CFR Part 160) and the OECD Good Laboratory Practice Regulations [C(97)186 Final]. The raw data and draft report will be audited by the WIL Quality Assurance Unit prior to submission to the Sponsor Representative to assure that the final report accurately describes the conduct and the findings of the study.

This study will be included on the

list of regulated studies.

# 11 RECORDS TO BE MAINTAINED:

All original raw data records, as defined by be stored as described in Section 12

#### 12 WORK PRODUCT:

The Sponsor will have title to all documentation records, raw data, slides, specimens and other work product generated during the performance of the study. Any remaining formulation and/or toxicokinetic samples will not be archived, but will be discarded after issuance of the final report. All work product, including raw paper

data, pertinent electronic storage media and specimens, will be retained for a period of 10 years following issuance of the final report in ti

monthly archiving fee for retention of all work product. All work product will be stored in compliance with regulatory requirements.

Any work product, including documents, specimens, and samples, that are required by this protocol. its amendments. or other written instructions of the Sponsor, to be

nackaged and labeled as defined by following delivery to the common carrier.

delivered to a common carrier 1 not be responsible for shipment

#### 13 REPORTS:

The final report will contain a summary, test article data, methods and procedures, appropriate individual animal and summary data tables, a copy of the protocol and amendments (if any) and an interpretation and discussion of the study results. The report will contain all information necessary to conform with current EPA specifications.

ide one (1) copy of an Audited Draft Report, submitted in a timely manner upon completion of the study prior to issuance of the final report. One (1) revision will be permitted as part of the cost of the study, from which Sponsor's reasonable revisions and suggestions will be incorporated into the Final Report, as appropriate. Additional changes or revisions may be made, at extra cost. It is expected that the Sponsor will review the draft report and provide comments a two (2) month time frame following submission. will submit the Final Report within one (1) month following receipt of comments. If the Sponsor's comments and/or authorization to finalize the report have not been received a n one year following submission of the draft report, elect to finalize the report following appropriate written notification to the Sponsor. Two (2) electronic copies of the Final Report on CD-R will be provided; requests for additional electronic or paper copies of the Final Report may result in additional charges.

# 14 ANIMAL WELFARE ACT COMPLIANCE:

This study will comply with all applicable sections of the Final Rules of the Animal Welfare Act (AWA) regulations (9 CFR Parts 1, 2 and 3). The Sponsor should make particular note of the following:

 The Sponsor Representative's signature on this protocol documents for the Study Director the Sponsor's assurance that the study described in this protocol does not unnecessarily duplicate previous experiments.

- Whenever possible, procedures used in this study have been designed to avoid or minimize discomfort, distress or pain to animals. All methods are described in this study protocol or in written laboratory standard operating procedures.
- Animals that experience severe or chronic pain or distress that cannot be relieved
  will be painlessly euthanized as deemed appropriate by the veterinary staff and
  Study Director. The Sponsor will be advised by the Study Director of all
  circumstances which could lead to this action in as timely a manner as possible.
- Methods of euthanasia used during this study are in conformance with the above-referenced regulation.
- The Sponsor/Study Director has considered alternatives to procedures that may
  cause more than momentary or slight pain or distress to the animals and has
  provided a written narrative description (AWA covered species) of the methods and
  sources used to determine that alternatives are not available.

#### 15 PROTOCOL MODIFICATION:

Modification of the protocol may be accomplished during the course of this investigation. However, no changes will be made in the study design without the verbal or written permission of the Sponsor. In the event that the Sponsor verbally requests or approves a change in the protocol, such changes will be made by